Prod. of the luffin-g gene. Homologous to chicken nov ge	finds:	en:	: 701	GENESEQP:R31608
! prod. of the luffin-g gene.	THUS:			GENESEKT . NASSTO
))	len: 278	ck: 9157	GENESEOD . 829910
- prod of	finds: 1	len: 277	ck: 429	GENESEQP:R29909
1 HCV NS4-NS5 peptide N29-3. New	finds: 1	len: 167	ck: 2886	GENESEQP: R29897
1 ! HCV NS4-NS5 peptide N29-2. New	finds: 1	len: 171	ck: 8862	GENESEQP: R29896
1 ! HCV NS4-NS5 peptide N29-1. New	finds: 1	len: 171	ck: 8862	GENESEQP: R29895
1 ! HCV NS4-NS5 peptide 2918. New	finds: 1	len: 285	ck: 8945	GENESEQP:R29879
l ! HCV NS4-NS5 peptide N29-1, N29	finds: 1	len: 171	ck: 9846	GENESEQP:R29874
HRSV 22K protein. Vaccines for	finds: 1	len: 194	ck: 9668	GENESEQP:R25303
! HK16. Hepatitis C virus antige	finds: 1	len: 253	ck: 4644	GENESEQP:R25891
! HCV polypeptide 16. Hepatitis	finds: 1	len: 239	ck: 3783	GENESEQP:R25869
. ! NANBH virus antigen. Antigenic	finds: 1	len: 85	ck: 5700	GENESEQP:R25601
<pre>! blaC31 mutation (5).</pre>	finds: 1	len: 286	ck: 4919	GENESEQP:R22695
! Protein transcribed from the	finds: 1	len: 297	ck: 7059	GENESEQP:R23004
NADH dehydrogenase subunit 2.	finds: 1	len: 482	ck: 617	GENESEQP:R21409
! VP6 inner capsid epitope, 84 T	finds: 1	len: 25	ck: 4995	GENESEQP: R20004
! HCV-T (6110-6557) encoded epit	finds: 1	len: 149	ck: 3078	GENESEQP:R14247
! Mouse kidney cell Band 3-1	finds: 1	len: 292	ck: 2381	GENESEQP: P60645
! Murine Cytotoxic Cell Protease	finds: 1	len: 247	ck: 528	GENESEQP:R13252
! Luffa cylindrica bioactive pro	finds: 1	len: 248	ck: 7673	GENESEOP: R12468
! Acetoacetyl CoA reductase enzy	finds: 1	len: 246	ck: 3836	GENESEQP:R10974
! Sequence of rhinovirus HRV2 vi	finds: 1	len: 323	ck: 5281	GENESEQP: P81093
! Sequence of rhinovirus HRV2 vi	finds: 1	len: 95	ck: 3060	
! Bovine RSV strain A 51908 M2 p	finds: 1	len: 186	ck: 5872	- 1
! Xenopus Bone Morphogenetic Fac	finds: 1	len: 398	ck: 1620	GENESEQP:R10995
! Regulatory protein VanS involv	finds: 1	len: 384	ck: 3600	
! Sequence encoding human respir	finds: 1	len: 194	ck: 9828	
! Sequence of human respiratory	finds: 1	len: 194	ck: 9828	
! Binding peptide 2 capable of b	finds: 1	len: 25	ck: 4995	
! Alpha-factor profibrolase from	finds: 1	len: 288	560	GENESEOP: P90460
! Acetyl-CoA reductase. Construd	finds: 1	len: 246	ck: 2829	GENESEQP: P94157
! Antigenic Epstein-Barr virus p	finds: 1	len: 48	ck: 718	GENESEQP: P91884 C
! Mouse gamma-2b chain New immun	finds: 1	len: 102	ck: 7203	GENESEQP: P90009 c

GENESEQP:R74180	GENESEQP:R72715	GENESEQP:R70100	GENESEQP:R71324	GENESEQP: R67596	GENESEQP: R66682	GENESEQP: R66680	GENESEQP: R66665	GENESEQP:R63160	GENESEQP:R63159	GENESEQP: R66355	GENESEQP:R62507	GENESEQP:R63444	GENESEQP:R63443	GENESEQP:R58587	GENESEQP:R58636	GENESEQP:R58635	GENESEQP:R63602 c	GENESEQP:R63592 c	GENESEQP:R58556 c	GENESEQP:R53748 c	GENESEQP:R53646 c	GENESEQP:R53251 c	GENESEQP:R50082 c	GENESEQP:R50081 c	GENESEQP:R49589 c	GENESEQP:R49438 C	GENESEQP:R49299 c		-	GENESEQP:R52667 cl	GENESEQP:R42317 cl	GENESEQP:R39260 ck:	GENESEQP:R39259 ck:	GENESEQP:R37294 ck:	GENESEQP:R32192 ck:
ck: 816	ck: 516	ck: 4652	ck: 7564	ck: 5465	ck: 5012	ck: 5136	ck: 4995	ck: 1814	ck: 2293	ck: 1046	ck: 2303	ck: 7811	ck: 8722	ck: 9708	ck: 3320	ck: 2685	ck: 5661	ck: 2807	ck: 4528	ck: 2863	ck: 4677	ck: 2035	ck: 108	ck: 3747	ck: 4300	ck: 4300	ck: 1278	ck: 3126	ck: 4901	ck: 5723	ck: 3489	k: 8110	k: 8652	c: 7673	c: 2829
3 len: 248	len: 359	len: 274	len: 329	len: 331	len: 25	len: 25	len: 25	len: 375	len: 376	len: 216	len: 313	len: 120	len: 120	len: 246	len: 124	len: 124	len: 367	len: 79	·len: 175	len: 355	len: 380	len: 56	len: 215	len: 352	len: 49	len: 49	len: 23	len: 24	len: 25	len: 303	len: 345	len: 391	len: 391	len: 248	len: 246
finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1	finds: 1
! Type I ribosome-inactivatin	! hisC gene product of Salmon	! Lettuce infectious yellows	! Acetyl-CoA-reductase. Trans	! A. aculeatus pectin methyle	Peptide MONOSER. Immunogeni	S	! Binding peptide B (84 TS) I	tiati	! Mouse growth differentiatio	! Ii protein. Identification	! Galactosyl transferase 3' c	! Amino terminal sequence of	! Amino terminal sequence of	! Nicotineamide adenine dinuc	! Amylase inhibitor protein 0		! MAP-kinase-phosphatase CL10	tase	! Heparin-binding secretory t	! Seven transmembrane recepto	! c-fos gene product. New ant	! Consensus sequence of signa	! NANBH virus antigenic fragm	! NANBH virus antigenic fragm	! Sequence of HLA-DR alpha ch	! Minigene HLA-DRalpha chain/	! Invarient chain (11) positi	! Invarient chain (11) positi	! Invarient chain (li) positi	! Equine herpesvirus US2 gene	! EBV VCA-p40 from ORF BdRF1.	Murine somatostatin recepto	! Human somatostatin receptor	! Plant type I RIP Luffin. An	by the a

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GENESEQP: W20776 GENESEQP:W20886 GENESEQP: W20571 GENESEQP:W06935 GENESEQP:W13272 GENESEQP: W06943 GENESEQP:W11479 GENESEQP:W11481 GENESEQP: W04831 GENESEQP: W01743 GENESEQP: R99253 GENESEQP:W00725 GENESEQP:R99316 GENESEQP:R83015 GENESEQP: R94586 GENESEQP: R94580 GENESEQP:R94579 GENESEQP: W00624 GENESEQP: R99576 GENESEQP: R99501 GENESEQP: R99695 GENESEQP:R91225 GENESEQP: R95055 GENESEQP: R95054 GENESEQP: R77433 GENESEQP: R89898 GENESEQP: R89897 GENESEQP: R89896 GENESEQP: R94990 GENESEQP: R94989 GENESEQP: R91274 GENESEQP: R75202 GENESEQP: R81333 GENESEQP: R81330 GENESEQP: R81327 GENESEQP: R81323 GENESEQP:R84103 č Č ck: čķ: č, ck: сķ: ck: č ck: ck: č. cķ: ck: ck: 5439 ck: ck: 8391 ck: 2766 ck: ĊĶ: ck: 2030 ck: 4995 ck: 7811 с С č ck: ck: ck: ck: ĊĶ: č čķ: č K ck: č, ck: 6185 2872 1679 6963 8017 9316 2018 9389 5195 9728 4145 1679 4323 9728 3773 2400 7433 9286 2940 7276 206 6122 5928 1456 5431 9019 9504 824 8013 2629 5723 len: 488 len: 276 154 136 140 114 244 221 205 207 259 374 207 371 458 259 271 358 83 244 25 328 421 160 342 457 437 423 282 354 62 225 142 123 140 123 303 finds: finds: 1 finds: 1 finds: 1 finds: 1 finds: finds: 1 finds: finds: finds: 1 finds: 1 finds: 1 finds: 1 finds: 1 finds: finds: 1 finds:] finds: 1 finds:] finds: 1 finds: finds: 1 finds: finds: finds: Ξ H. pylori flagella-associated H. pylori secreted or CagI locus product 6. Helicob CagI locus product 14. Helico D. immitis venom allergen ant Vascular endothelial growth fa Rhodococcus erythropolis SK92 D. immitis mature venom aller C. pneumoniae 53 kDa antigen. Cytoplasmic antiproteinase-2 C. pneumoniae polypeptide ant: Vascular endothelial growth fa Human thyroid transcription Human SH-PTP1 variant derived C. pneumoniae polypeptide ant: Chlamydia pneumoniae polypept SacI endonuclease. DNA encodi Wasp venom BrhTX-1 subunit (b VP6 binding peptide B. Encaps Nitrilase regulatory factor Human placenta G-protein coup IL-2-DETA-DGAL4 multidomain pr TGF-a-DETA-DGAL4 multidomain Antigen of Rochalimaea hensel Rat kynurenine aminotransfera Rat kynurenine aminotransfera Rat kynurenine aminotransfera Nsk2 receptor tyrosine-kinase Nsk2 receptor intracellular Eimeria Tyrosine phosphatase MPTP-del Human VLA-4 reshaped antibody Humanized VLA-4 antibody 21.6 Mouse anti-VLA-4 antibody 21 Mouse VLA-4 antibody 21.6 ligh Equine herpesvirus (EHV1) unic pylori secreted or gametocyte antigen. Pr peripla

GENESEQP: W28052 GENESEQP: W51011 GENESEQP:W37816 GENESEQP:W56137 GENESEQP: W55815 GENESEQP: W55590 GENESEQP: W50006 GENESEQP: W47606 GENESEQP:W34214 GENESEQP:W41747 GENESEQP: W38190 GENESEQP: W45093 GENESEQP:R63906 GENESEQP:W29525 GENESEQP: W29524 GENESEQP: W29523 GENESEQP: W24493 GENESEQP: W24492 GENESEQP: W24491 GENESEQP: W34668 GENESEQP:W17971 GENESEQP:W26413 GENESEQP:W22428 GENESEQP:W22413 GENESEQP:W22410 GENESEQP:W25146 GENESEQP:W25141 GENESEQP:W13647 GENESEQP:W10031 GENESEQP: W21709 GENESEQP:W21704 GENESEQP:W09433 GENESEQP: W20139 GENESEQP: W20696 GENESEQP: W20324 GENESEQP:W20259 č Č č; ck: 7216 ck: 5054 č Č öķ: ck: ck: 7408 ck: 428 ck: ck: 5054 ck: 1840 č, ck: ck: ck: 9381 ck: 7673 ck: 1423 č; ς; č. čķ: Š ck: ck: ck: <u>с</u> cķ: č ck: ck: cy. č, č; ck: 1367 ck: 5141 3813 2273 5567 9828 3783 3320 2685 3320 2685 1423 9097 6033 3737 9019 9557 2629 7681 7673 5362 8373 7681 7673 8653 7893 6372 len: 140 len: len: len: len: len: len: len: len: len: 307 365 194 107 176 239 271 248 124 122 124 124 122 124 438 47 142 121 123 250 248 30 74 250 248 328 102 108 53 121 finds: 1 finds: finds: 1 finds: finds: 1 finds: finds: finds: finds: finds: finds: finds: finds: 1 finds: 1 finds: 1 finds: 1 finds: finds: finds: finds: finds: finds: finds: finds: finds: 1 finds: 1 Human secreted apoptosis-re Amino acid sequence of phos H. pylori ORF 09cp10224_429 Human liver carbonyl reduct Streptomyces roseofulvus fr Human papillomavirus-18 E2 HRSV protein 22K. Productio Hepatitis C virus antigen. Streptomyces ketoreductase Type I ribosome-inactivatin Wheat amylase inhibitor 0.2 Ξ. Open reading frame 2 Soybean SCARECROW SRPg1 pro Residues 217-392 of human Wheat amylase inhibitor 0.1 Wheat amylase inhibitor 0.2 Novel amylase inhibitor pro Novel amylase inhibitor pro Novel amylase inhibitor pro Arabidopsis thaliana Rec-A RAC protein kinase C-termin Swinepox virus HindIII C Humanised alpha-4 integrin Alpha-4 integrin mouse MAb Luffin-B (a ribosome inhibi Humanised alpha-4 integrin Luffin-A (a ribosome inhibi Invariant chain region whic Protein encoded by clone Luffin-B. Inactive precurso Luffin-A. Inactive precurso Human placenta purinergic p H. pylori cytoplasmic H. pylori secreted or perip Ξ. pylori cell envelope tra pylori secreted or perip peptid

Ë	! A Staphylococcus aureus proce	inds: 1	fin	1: 67) len:	: 8790	2 ck:	. GENESEQP: w79302
. <u> </u>	partyl t	inds: 1	н	161	len:	: 3773	ck:	GENESEOP: W38716
	protei	inds: 1	ь	: 100	len:	5825	ck:	GENESEQP: W77666
	Ω,	finds: 1		: 212	len:	4556	ck:	GENESEQP: W70962
هــــه	neumoniae pi	inds: 1	н	: 174	len:	5404	ck:	GENESEQP:W38580
		inds: 1	н	: 120	len:	3575	Ċ.	GENESEQP:W64373
	- 0	inds: 1	+	: 375	len:	1814	ck:	GENESEQP:W65460
<u> </u>	! Clone HTPEF86 of TM4SF superta	inds: 1	H	: 250	len:	1162	ck:	GENESEQP:W61619
	SOCS1 prot	ds: 1	finds:	: 211	len:	758	ck:	GENESEQP:W62616
	protein.	is: 1	finds	225	len:	9828	ck:	GENESEQP:W62615
		1s: 1	finds	: 198	len:	1060	ck:	GENESEQP:W62614
	protein.	ls: 1	finds	212	len:	4527	ck:	GENESEQP:W62613
	3 protein.	ls: 1	finds	134	len:	8778	ck:	GENESEQP:W62624
	protein.	S: 1	finds	207	len:	6307	ck:	GENESEQP:W62623
		s: 1	finds	130	len:	4000	ck:	GENESEQP:W62622
	OCS1 prot	s: ⊢	finds:	212	len:	5610	ck:	GENESEQP:W62617
	FCR-II protein sequence. Nucle	s: 1	finds:	263	len:	3029	ck:	GENESEOP:W69232
	_	s: 1	finds	387	len:	843	ck:	GENESEQP: W30679
		s: 1	finds:	369	len:	8912	ck:	GENESEQP:W62612
		5:	finds:	61	len:	8918	ck:	GENESEQP:W44882
	beta-turn	1-	finds	104	len:	9655	ck:	GENESEQP:W44879
	beta-turn	:: -	finds	167	len:	1990	ck:	GENESEQP:W44910
	turn	بر ۳	finds:	110	len:	4699	ck:	GENESEQP:W44909
	beta-turn	:: 	finds:	79	len:	1205	ck:	GENESEQP:W44908
		н	finds:	127	len:	3740	ck:	GENESEQP:W44907
	beta-turn	 1	finds:	70	len:	6449	ck:	GENESEQP: W44906
	beta-turn	 L	finds:	39	len:	1135	ck:	GENESEQP:W44905
	TD.	 ,	finds:	271	len:	5290	ck:	GENESEQP:W57889
	interleukin l	 1	finds:	260	len:	7422	ck: 7	GENESEQP:W60764
	eumoniae DHAT	 H	finds:	387	len:	843	ck: e	GENESEQP:W60267
	glyce		finds:	339	len:	4443	ck: 4	
		 P	finds:	24	len:	3126	ck: 3	
	Human V28 seven transmembrane	 L	finds:	355	len:	2863	ck: 2	
	Sunflower antifungal protein M	<u>-</u>	finds:	118	len:	4306	ck: 4	
	! Flea serine protease inhibitor	<u></u>	finds:	137	len:	7060	ck: 7	
	Glycerol-3-phosphate dehydroge	ь	finds:	339	len: 3	4443	ck: 4	_
	- 7	ь	finds:	71	len: 7	9911 :	ck: 9	GENESEQP:W59024

! H. pylori GHPO 1099 procein	finds: 1	len: 115	ck: 2117	GENESEQP:W98226
1 GHPO 1075	finds: 1	len: 102	ck: 3203	GENESEQP: W98755
GHPO 16/0	finds: 1	len: 304	ck: 4789	GENESEQP:W98862
ortierella alpina u	finds: 1	len: 446	ck: 5663	GENESEQP: W95506
tis ele	finds: 1	len: 161	ck: 9212	GENESEQP:W83391
in encode	finds: 1	len: 221	ck: 9823	GENESEQP: W88568
de fragment ence	finds: 1	len: 36	ck: 1342	GENESEQP:W89100
with 95%	finds: 1	len: 351	ck: 8607	GENESEQP: W85591
with 98%	finds: 1	len: 354	ck: 2306	GENESEQP:W85590
rom cluste	finds: 1	len: 343	ck: 8829	GENESEQP:W89910
റ	finds: 1	len: 343	ck: 9038	GENESEQP: W90011
! Human ATG-1709 protein. New	finds: 1	len: 317	ck: 3945	GENESEQP:W73507
curase enz	finds: 1	len: 446	ck: 5663	GENESEQP: W85119
-2 truncated	finds: 1	len: 163	ck: 3123	GENESEQP:W86217
	finds: 1	len: 168	ck: 2203	GENESEQP:W86216
	finds: 1	len: 173	ck: 4363	GENESEQP:W86215
	finds: 1	len: 178	ck: 5797	GENESEQP:W86214
(VEGF-relate	finds: 1	len: 185	ck: 8227	GENESEQP:W86235
! Amino acid sequence of the	finds: 1	len: 420	ck: 8848	GENESEQP:W68541
fact	finds: 1	len: 206	ck: 9273	
s II pep	finds: 1	len: 24	ck: 3126	
endot	finds: 1	len: 207	ck: 1679	
	finds: 1	len: 226	ck: 5027	
M. tuberculosis immunogenic	finds: 1	len: 120	ck: 3575	GENESEQP:W81740
! Human EPRG1 protein #2. New	finds: 1	len: 157	ck: 9756	
protein #1.	finds: 1	len: 225	ck: 447	
avirus protei	finds: 1	len: 368	ck: 9982	GENESEQP:W68494 c
S pneumoniae GTP binding P	finds: 1	len: 189	ck: 4504	GENESEQP:W80624 C
trile hydrau	finds: 1	len: 244	ck: 2018]	_
Bovine growth differentiate	finds: 1	len: 375	ck: 9305 1	GENESEQP:W69887 c
growth differentiat	finds: 1	len: 375	ck: 1463 l	
! Human growth differentiatio	finds: 1	len: 375	ck: 1814 l	
! Murine growth differentiati	finds: 1	len: 376	ck: 2293 l	
! Ovine growth differentiation	finds: 1	len: 375	ck: 1229 l	
	finds: 1	len: 375	ck: 1805 l	
! Rat growth differentiation	finds: 1	len: 376	ck: 1496 l	GENESEQP:W69889 cl

(178)(*)(*)**(*)**(*)**(*)**(*)**(*)**(*	A, P)XX(L, I, V, M, A, F, Y, W) (C, T, S) (R, K, H	mismatches T,S)(L,I,V,M,	<pre>RNS on geneseqp:* allowing ((X){1,200}(L,I,V,M,A,P)X(P, </pre>
R10995 ck: 1620 len: 398 Xenopus Bone Morphogenetic Factor BMP-2A			Total finds: 245 Total length: 23,686,106 Total sequences: 188,963 Total sequences: 188,963
1: LVIKLKKKKNDYSKLERKLYMYTVASIVYVLYIKSMIRGKLGDWILSILENKYDLNHLD SLLDEAPDMPVDQKAKYVHITLDKAYRLEQLIDEFFEITRYNLQTITLIKTHIDLY YMLVQMTDEFYPQLSAHGKQAVIHAPEDLTVSGDPDKLARVFNNILKNAAAYSEDNSIIDITAGLSGDVVSIEFKNTGSIPKDKL	Formatted on 9Jul1999	on 8Jul1999,	base
1	! Fragment of mouse hepatic STAT	1 finds: 1	GENESEQP:Y07241 ck: 3064 len: 141
	! Fragment of human hepatic STAT	1 finds: 1	GENESEQP:Y07240 ck: 2614 len: 141
TT	! Fragment of human placental ST	1 finds: 1	GENESEQP:Y07239 ck: 2144 len: 141
1. MCDRUCKEFIRGHCINGEPCHRCHURFEUDDIALIUDONFRINGTERACIONT CETICO	! Mycobacterium species protein	0 finds: 1	GENESEQP:Y04743 ck: 5429 len: 110
LIVION CY. 2000 TEH. 124 : Seduence encounting intimate reconstruction	! Mycobacterium species protein	finds: 1	GENESEQP:Y04901 ck: 6286 len: 72
of 0828 les. 104 Secuence encoding human regularators	! Mycobacterium species protein	0 finds: 1	GENESEQP:Y04960 ck: 8320 len: 280
I: MSKKNECKFEIRGHCLNGKKCHFSHNIFEWFHALLVKQNFMLNKILKSMUKSIUILSEISGA ESTVSDINDHAKNNDII	! Mycobacterium species protein	3 finds: 1	GENESEQP:Y04959 ck: 8454 len: 273
(A) [1, AUU] [1, 1, V, M, A, F) A(F) 1, B) [1, 1, V, M, A, F, X, V, M, A, X,	! Human regulatory molecule HRM-	2 finds: 1	GENESEQP:W93953 ck: 7759 len: 352
"\ C# \ \	! Breast cancer associated antig	7 finds: 1	GENESEQP:Y07007 ck: 4692 len: 177
15. 0000 los. 104 l gomono of hims topping	! Breast cancer associated antig	7 finds: 1	GENESEQP:Y07041 ck: 1679 len: 207
1: CNIAPASIVSRNIVYTRAQPNQDIA	! Wheat Type III glutathione tra	3 finds: 1	GENESEQP:Y05545 ck: 5919 len: 243
$1 < (X) \{1, 200\}(L, I, V, M, A, P)X(P, T, S)(L, I, V, M, A, P)XX(L, I, V, M, A, F, Y, W, Y, Y,$! Human 5' EST secreted protein	2 finds: 1	GENESEQP:Y11993 ck: 4119 len: 102
P81999 ck: 4995 len: 25 Binding peptide 2 capable of binding to	! Human CLIP immunomodulatory pe	finds: 1	GENESEQP:Y05204 ck: 3125 len: 24
LGMNHUGNQCHUGANSU YMAAMESUQFSALE SUCSAAU I Q'E ET YNNEQC LENAF	! Brn-3a polypeptide. New polype	3 finds: 1	GENESEQP:W96262 ck: 3324 len: 423
I: MKEESTETAVLEAASSALAAPVANTTEEETTAVLEAASSALAAVANTOTELEHA HDNAQLLTAIDEDGDTVGLAYVGGMCQLKHSTGVIDHSAINLLVALTMAHELGHN TONNINGANATATATATATATATATATATATATATATATATATAT	! Human 5' EST secreted protein	3 finds: 1	GENESEQP:Y11970 ck: 1817 len: 113
1. \(\begin{array}{c} \\\ \begin{array}{c} \(\begin{array}{c} \\\ \begin{array}{c} \(\begin{array}{c} \\\ \begin{array}{c} \(\begin{array}{c} \\\ \begin{array}{c} \\ \begin{array}{c} \\\ \begin{array}{c} \\\ \begin{array}{c} \\\ \begin{array}{c} \\\ \begin{array}{c} \\\ \begin{array}{c} \\\ \	! Fragment of human secreted pro	finds: 1	GENESEQP:Y02934 ck: 2639 len: 53
, o	! S. aureus polypeptide. New ess	5 finds: 1	GENESEQP:Y03782 ck: 881 len: 135
	! S. aureus polypeptide. New ess	4 finds: 1	GENESEQP:Y03794 ck: 7277 len: 144
VNIVSPOITATUMVAAIKQUVUUKIVATTPVAKLGUPEETASICAMUSSEESGES! GADFSINGGIHMG	! Winged bean chymotrypsin inhib	7 finds: 1	GENESEQP:W97840 ck: 7697 len: 207
1: XTECHNICAL X (2) (L) XX (2) (L) XX (1) XXXX (A) X (1118) MTGRIAYTGGMGGIGTAICQRIAKDGFRVVAGCGPNSPRREKWLEQQKALGFDFIASEGNVA	! Murine myostatin. Increasing m	6 finds: 1	GENESEQP:W97886 ck: 2420 len: 376
1 < (x)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,A,P,Y,W,P,Y,W,A,P,P,Y,W,P,P,Y,P,Y,P,Y,P,Y,P,Y,P,Y,P,Y,P	! Bovine myostatin (mutant form)	6 finds: 1	GENESEQP:W97885 ck: 1462 len: 286
P94157 ck: 2829 len: 246 ! Acetyl-CoA reductase. Constructing new p	! Bovine myostatin. Increasing m	5 finds: 1	GENESEQP:W97884 ck: 8371 len: 375
TO NOME CEDUCE DEFECTION I EFFECTANTARY FOR A PARTICULAR	! Human myostatin. Increasing mu	5 finds: 1	GENESEQP:W97887 ck: 1814 len: 375
$\begin{array}{cccc} & & & & & & & & & & & & & & & & & $! H. pylori ORF 06cp30603_107440	6 finds: 1	GENESEQP:Y11062 ck: 4555 len: 116
ryloo4 CK: /10 Ien: 40 ! Antigenic Epstein-Bair virus peptio	! H. pylori ORF hp6p22217_235640	2 finds: 1	GENESEQP:Y11009 ck: 2840 len: 142
	! Staphylococcus aureus mutant F	2 finds: 1	GENESEQP:W97722 ck: 1929 len: 302
1: FQVYILPPPAEQLSRKDVSLTCLVVGFNPGDISVEWTSNGHTEENYKDTAPVLDSDGSYFI	! Rhodococcus corallina ohp oper	4 finds: 1	GENESEQP:w99185 ck: 9793 len: 314
1 <(x)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W	! Secreted protein encoded by ge	finds: 1	GENESEQP:Y01440 ck: 9731 len: 64
p90009 ck: 7203 len: 102 ! Mouse gamma-2b chain New immunoglobulin	! Mannosyl transferase involved	6 finds: 1	GENESEQP:W88330 ck: 7734 len: 336

P90009	ck: 7203 len: 102 ! Mouse gamma-2b chain New immunoglobulin
!	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W X{6}(P)X(P)(A)XX(L)(S)(B)XX(Y)XXX(C)X(80) PQVYILPPPAEQLSRKDVSLTCLVVGFNPGDISVEWTSNGHTEENYKDTAPVLDSDGSYFI</pre>
P91884	ck: 718 len: 48 ! Antigenic Epstein-Barr virus peptide Ide
	<(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,WX(19)(P)X(P)(L)xx(A)(S)(R)xx(P)xxx(S)x(13) RGWFCPSLCPSEEPGTSGTPEPLGPASRRPPGLRSPLSPVKPKECLRG
P94157	ck: 2829 len: 246 ! Acetyl-CoA reductase. Constructing new p
1: VNTVSPGYIATDMVKAIRQDVI	1 <pre><(x){1,200}(L,1,V,M,A,P)X(P,T,S)(L,1,V,M,A,P)XX(L,1,V,M,A,F,Y,W</pre>
P90460	ck: 560 len: 288 ! Alpha-factor profibrolase from Southern
1: 1: HDNAQLLTAIDFDGDTYGLAYY	1 < (X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,X,P)XX(L,I,V,M,A,F,Y,W,X,P)XX(L,I,V,M,A,F,Y,W,X,P)XX(L)XXX(G)X{123}
P81999	ck: 4995 len: 25 ! Binding peptide 2 capable of binding to
: :	<(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{2}(I)x(P)(A)xx(V)(S)(R)xx(V)xxx(A)x{7} CNIAPASIVSRNIVYTRAQPNQDIA
P70477	ck: 9828 len: 194 Sequence of human respiratory syncytial
1: 1: ESTVSDINDHAKNNDTI	<pre><(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{83}(I)x(S)(I)xx(I)(T)(K)xx(A)xxx(M)x{95} wsrnpckfeirghclngkrcheshnyfewpphallvrqnfmlnrilksmdksidtlseisga</pre>
P70785	ck: 9828 len: 194 ! Sequence encoding human respiratory sync
1: 1: ESTVSDINDHAKNNDTI	<pre><(x){1,200}(L,I,V,M,A,F)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{83}(I)x(S)(I)xx(I)(T)(K)xx(A)xxx(M)x{95} MSRRNPCKFEIRGHCLNGKRCHFSHNYFEWPPHALLVRQNFMLNRILKSMDKSIDTLSEISGA</pre>
R24296	ck: 3600 len: 384 ! Regulatory protein VanS involved in glyc
1: 1: 1: 1: 1: 1: 1: 1: 1: 1: 1: 1: 1: 1	<pre><(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{84}(V)x(S)(I)xx(L)(S)(R)xx(L)xx(A)x(284) LVIKLKNKKNDYSKLERKLYMYIVATVYVYLTRSMIRGKLGDWILSILENKYDLNHLD LYKLKNKKNDYSKLERKLYMYIVATVYVYLTRSMIRGKLGDWILSILENKYDLNHLD LYKAYRLEQLIDEFEITRYNLQTITLTKTHIDLY</pre>
(MLVQMTDEFYPQLSAHGKQA)	IMLVQMTDEFYPQLSAHGKQAVIHAPEDLTVSGDPDKLARVFNNILKNAAAYSEDNSIIDITAGLSGDVVSIEFKNTGSIPKDKL

R29879 CK: 8945 len: 285 ! HCV NS4-NS5 peptide 2918. New hepatitis <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)X(L,I,V,M,A,F,Y,W x{18}(L)x(S)(V)xx(A)(C)(K)xx(P)xxx(A)x{251} x{18}(L)x(S)VXX(A)(C)(K)xx(P)xxX(A)X(251) rrdvlkdmkakastvkakllsvebackltpphsakskfdrgakdvgslsskavnhihsvmkdl 1: vrdvlkdmkakastvkakllsvebackltpphsakskfdrgakdvgslsskavnhihsvmkdl	R20004 ck: 4995 len: 25 ! VP6 inner capsid epitope, 84 TS, Peptide B. <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C,X){1,2}(I)X(P)(A)xx(V)(S)(R)xx(V)xxx(A)x{7} 1: CNIAPASIVSRNIVYTRAQPNODIA	н
R29874 ck: 9846 len: 171 ! HCV NS4-NS5 peptide N29-1, N29-3 <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W) <(X)[1,200](L,X(S)(V)XX(A)(C)(K)XX(P)XXX(A)X(137) X(18](L)X(S)(V)XX(A)(C)(K)XX(P)XXX(A)X(137) YRDVLKEMKAKASTVKAKLLSVEEACKLTPPHSARSKFGYGAKDVRSLSSKAVNHIRSVWXDL	R14247 ck: 3078 len: 149 ! HCV-T (6110-6557) encoded epitope. New nucl <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C <(X){1,200}(V)XX(A)(C)(K)XX(P)XXX(A)X(118) X{15}(L)X(S)(V)XX(A)(C)(K)XX(P)XXX(A)X(118) VLKBSKAKASTVKAKLLSVEBACKLTPPHSARSKFGYGAKDVRSLSSRAINHIRSVWKDLLEDTET 1:	ь
TIEC A', A	P60645 ck: 2381 len: 292 ! Mouse kidney cell Band 3-like anion transport of the control of the c	1 VTSLNGIOI TAASLAFPI
R25891 ck: 4644 len: 253 ! HK16. Hepatitis C virus antigen expresse <(X)[1,200](L,I,V,M,A,P)X(P,T,V,M,A,P)XX(L,I,V,M,A,F,Y,W 1	A vect	1 KNRYNKTNQ SSFLSWIKK
R25869 ck: 3783 len: 239 ! HCV polypeptide 16. Hepatitis C virus an <(x)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(8)(L)X(S)(I)XX(A)(C)(K)XX(P)XX(A)X(139) VCSMSYTMCAALTECAALESKLPINPLSNSLLRHHSMYYSTISRSASLRQKKVTFDRLQVL LYDVVSTLPQAVMGPSYGFQYSPGQRVEFLVNTWKSKKCPMGFSYDTRCFDSTVTE NDIRTE	R12468 ck: 7673 len: 248 ! Luffa cylindrica bioactive protein. New protein	1 QVPSLATIS! TKNIQLLLN'
R25601 ck: 5700 len: 85 ! NANBH virus antigen. Antigenic polypepti <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(32](L)x(S)(V)xx(A)(C)(K)xx(P)xxx(A)x(37) KKVTFDRLQVLDDHYRDVLKEMKAKASTVKAKLLSVEBACKLTPPHSAKSKFGYGAKDVRN 1: KKVTFDRLQVLDDHYRDVLKEMKAKASTVKAKLLSVEBACKLTPPHSAKSKFGYGAKDVRN	2 (X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) (X)(1)(X)(X)(X)(X)(X)(X)(X)(X)(X)(X)(X)(X)(X)	1 VNTVSPGYIA GADFSLNGGI
R22695 ck: 4919 len: 286 ! blaC31 mutation (5). Novel DNA modificat (X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W (1)3)(A)X(S)(V)XX(A)(S)(R)XX(L)XXX(M)X(77) MSTOCHEWALLEPFFAAFCLPVEAHPETLVKVKDAEDQLGARVGYIELDLNSGKILESFRPEER TMPAAMATTLRKLLTASSVTLASRQQLRDWMEADKVAGPLLRSALPAGWFIADKS GAGERGSRGIIAALGPDGKPSRIVVIYTTGSQATMDERNRQIAEIGASLIKHW GAGERGSRGIIAALGPDGKPSRIVVIYTTGSQATMDERNRQIAEIGASLIKHW	<pre>1</pre>	1 GDDMTLFCQM RRFFLDLDII
R23004 ck: 7059 len: 297 ! Protein transcribed from the mba sequenc (x){1,200}(L,T,V,M,A,P)X(P,T,S)(L,T,V,M,A,P)X(L,I,V,M,A,F,Y,W)	P81092 ck: 3060 len: 95 ! Sequence of rhinovirus HRV2 viral protein F <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C ×(27)(I)X(P)(V)XX(I)(S)(K)XX(I)XX(I)X{52} 1: GVTDYIHMLGEAFGNGFVDSVKEHIHAINPVGNISKKIIKWMLRIISAMVIIIRNSSDPQTILA P81093 ck: 5281 len: 323 ! Sequence of rhinovirus HRV2 viral protein F	н
R21409 ck: 617 len: 482 i NADH dehydrogenase subunit 2. New DNA se <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)X(L,I,V,M,A,F,Y,W x(184](L)x(S)(L)xx(F)(S)(K)xx(L)xxx(M)x(282) x(184)(L)x(S)(L)xx(F)(S)(K)xx(L)xxx(M)x(282) x(184)(L)x(S)(L)xx(F)(S)(K)xx(L)xx(M)x(282) x(184)(L)x(S)(L)xx(F)(S)(K)xx(L)xx(M)x(282) x(184)(L)x(S)(L)xx(F)(S)(L)xx(F)(X)x(L)xx(M)x(B)X(L)X(L)X(B)X(B)X(B)X(B)X(B)X(B)X(B)X(B)X(B)X(B	R24190 ck: 5872 len: 186 ! Bovine RSV strain A 51908 M2 protein. Bovin <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x(83)(L)X(S)(I)XX(I)(T)(K)XX(A)XXX(M)X{87} MSRRNPCKYEIRGHCLNGKKCHFSHNYFEWAPHALLVRQNFMLNKILKSMDRNNDTLSEISGAAEL 1:	1 DINVDEQNE

1 <(x)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x(37)(L)X(S)(A)XX(A)(S)(R)XX(L)XXX(S)X(195) 1: DVRESLSGSSSTSYSETGDLKRALPSNGTYYNLTILLSSASGASRYTLMTLSNYDGKAITVAVDV QVPSLATISLENSLWSALSKQIQLAQTNNGTFKTPVVITDDKGQRVEITNVTSKVV	R37294 ck: 7673 len: 248 Plant type I RIP Luffin. Analogues of type	2(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C,X(112)(L)x(S)(L)x(Y)(T)(K)xx(I)xxx(A)x(A)x(118) 1:	1 <(x)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x[54](P)x(P)(I)xx(I)(T)(K)xx(L)xxx(G)x{36} 1: LLTLSPYXKVFLARLIWWLQYFITRAEAHLQVWIPPLNVRGGRDAIILLACAVHPEPIFDITKX R32192 ck: 2829 len: 246 ! Sequence encoded by the acetoacetyl-CoA red	<pre>1</pre>	R31608 ck: 7012 len: 205 ! Homologous to chicken nov gene exon 3-4-end	R29910 ck: 9157 len: 278 ! Prod. of the luffin-g gene. New gene coding <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C	R29909 ck: 429 len: 277 ! Prod. of the luffin-f gene. New gene coding 1 <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x(57)(L)x(S)(A)xx(A)(S)(R)xx(L)xxx(S)x(204) 1: MKRFTYLILAITVAASTVEADVRESLSGSSSTSYSKFIGDLRKALPSNGTVYNITLLLSSASGASH AEASRFKYIEGQIIERISKNQVPSLATISLENEWSALKYOLAQTNNGTFKTPVV ITDDKGGRVEITNYTSKVVTKNIOLLLNYKGNVAAFDEDVOAKH	1 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C)(K)XX(P)XXX(A)X{133} 1: YRDVLKEMKAKASTVKAKLLSVEEACKLTPPHSARSKFGYGAKDVRSLSSKAVNHIRSVWKDLLED	R29896 ck: 8862 len: 171 ! HCV NS4-NS5 peptide N29-2. New hepatitis C (X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x(18)(L)x(S)(Y)xx(A)(C)(K)xx(P)xxx(A)x(137) 1: YRDVLKEMKAKASTVKAKLLSVEEACKLTPPHSARSKFGYGAKDVRSLSSKAVNHIRSVWKDLLED R29897 ck: 2886 len: 167 ! HCV NS4-NS5 peptide N29-3. New hepatitis C	R29895 CK: 8862 len: 171 ! HCV NS4-NS5 peptide N29-1. New hepatitis C <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x(18](L)x(S)(V)xx(A)(C)(K)xx(P)xxx(A)X(137] 1: YRDVLKEMKAKASTVKAKLLSVEEACKLTPPHSARSKFGYGAKDVRSLSSKAVNHIRSVWKDLLEG	QCCDLDPEARQAIRSLTERLYIGGPLTNSKGQNCGYRRCRASGVLTTSCGNTLTCY LKASAACRAAKLQDCTMLVCGDDLVVICESAGTQEDAANLRVFTEAMTRNSA
1 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(26){P}x(P)(P)xx(V)(S)(K)xx(M)xxx(L)x{7} 1: MAISGYPVLGFFIIAVLMSAQESWALPKPPKPVSKMRMATPLLMQALPM	R49589 ck: 4300 len: 49 ! Sequence of HLA-DR alpha chain leader pe	R49438 ck: 4300 len: 49 ! Minigene HLA-DRalpha chain/Ii 24 residue (X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(26}(P)x(P)x(P)(S)(K)xx(L)xx(L)x(I) 1: wAISGVFVLGFFIIAVLMSAQESWALPKPPKPVSKNRMATFLLMQALPM	R49299 ck: 1278 len: 23 ! Invarient chain (11) position 97-119. No <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(P)x(P)(P)xx(V)(S)(K)xx(M)xxx(L)x(6) 1: LPKPPKPVSKMRMATPLLMQALP	R49297 ck: 3126 len: 24 ! Invarient chain (li) position 97-120. No <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(P)x(P)(P)xx(V)(S)(K)xx(M)xxx(L)x{7} 1: LPKPPKPVSKMRMATPLLMQALPM	1: <(X){1,200}(L,1,V,M,A,P)X(P,T,S)(L,1,V,M,A,P)XX(L,1,V,M,A,F,Y,W,X(P)x(P)x(P)(P)xx(V)(S)(K)xx(M)xxx(L)x(8)	<pre><(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,X,X,E)(Z,E)</pre>	**(51)(A) X (F) (A) X (F) (A) (A) X (F) (A) (A) X (F) (A) (A) (A) (A) (A) (A) (A) (A) (A) (A	R42317 ck: 3489 len: 345 EBV VCA-p40 from ORF BdRF1. Peptide(s) a	R39260 ck: 8110 len: 391 ! Murine somatostatin receptor-1. Somatost (X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(188){1}x(P)(I)xx(F)(S)(R)xx(A)xxx(G)x[187) x(188)(I)xx(P)(S)(R)xx(A)xxx(G)x[187) yerpograssessessessessessessessessessessessesse	R39259 ck: 8652 len: 391 ! Human somatostatin receptor-1. Somatosta (X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(188){I)x(P)(I)xx(F)(S)(R)xx(A)xxx(G)x(187) MEPNOTASSPSSSSSSSCGEGGGSRCPGAGAADGMEEPGRNASONGTLSEGQGSAILISE NLGVWVLSLLVILDIVVFSRTAANSDGTVACMMLAVEPAQAWLVGFVLYTTFLMGFL LPVGAICLCYVLIIAKMRMVALKAGMQQRKRSERKITLMVMMVVMVFVICWMPFTVVVQLVNVFAEQDDATVSQLSVILGYANSCA	TKNIQLLLNYKQNVA

. .

R66680 ck: 5136 len: 25 ! Peptide 84 TS-CYS. Immunogenic complex c	Ì
1 <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{2}(I)x(P)(A)xx(V)(S)(R)xx(V)xxx(A)x[7] 1: CNIAPASIVSRNIVYTRAQPNQDIA	R58635 ck: 2685 len: 124 Amylase inhibitor protein 0.26 AI. New amylase inhibitor protein 0.26
R66665 ck: 4995 len: 25 ! Binding peptide B (84 TS) Immunogenic co	SSNEMBAIDEIDSIKNAGGRVFVHCQAGISRSATICLAYLMRTNRVKLDEAFEFVKQRRSIISBNESEMGQLLQFESUVLFHSUSAEL
x(65)(L)x(T)(A)x(L)(S)(K)xx(L)xx(L)xx(L)xx(L)xx(L)xx(L)xx(L)x	1 <-(X){1,200}(L,I,Y,M,A,P)X(P,T,S)(L,I,Y,M,A,P)XX(L,I,Y,M,A,F,Y,W)(C) x(182)(L)x(S)(A)xx(A)(S)(R)xx(M)xxx(L)x(154) MYMEYGTLDAGGLEALLGERAAGCLLLDCRSFFAFNAGHIAGSYNVRFSTIVRRRAKGAMGLEHIV 1 DET VIGSAYHASRKOMIDALGITALINGSANCYMFEGHYQYKSIPVEDNHKADI
R63160 ck: 1814 len: 375 i Human growth dilleteiniación for servición (x) {1,200} (L,I,V,M,A,F)X(F,T,S)(L,I,V,M,P)XX(L,I,V,M,A,F,Y,W)	63602
IKPMADGIKKITGIASHAHAMSFOJGIAGA KATALAMSTOGLDCDEHSTESRCCRYPLIVDFEAFGWDWIIAPKRYKANYCSGECEFVF NGHDLAVTFPGPGEDGLNPFLEVKVTDTPKRSRRDFGLDCDEHSTESRCCRYPLIVDFEAFGWDWIIAPKRYKANYCSGECEFVF	1 $<(X)\{1,200\}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q,X,Y,Z,Y,Z,Y,Z,Y,Z,Y,Z,Y,Z,Y,Z,Y,Z,Y,Z,Y$
1 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,1,V,M,A,E,1,V,M,E,1,	:63592 ck: 2807 len: 79 MAP-kinase-phosphatase CL100 partial se
R63159 ck: 2293 len: 376 Mouse growth differentiation factor-8 pr	1 <(x)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q,Y,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,
1: MDDQNDLISNNEQLEMERAPGAPÉSKCSRGALYTGFSILVTLLLAGQATTAYFLYQQQGRLD SRHSLEQKPTDAPPKESLELEDPSSGLGVTKQDLGPVPM	R58556 ck: 4528 len: 175 ! Heparin-binding secretory transforming fact
1 <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{81}(P)x(P)(P)xx(V)(S)(K)xx(M)xxx(L)x{119}	DYPEVLOEIWPVLRNVEINFLGFLLPLLIMSYCYF KALUTLE SCARBAAAAAA LLVVIVFFLFWTPYNVMIFLETLKLYDFFPSCDMRKDLRLALSVTETVAFSHCCLNPLIYAFAGEKFRRYLYHLYGKCLAVLCGRSVH
R66355 ck: 1046 len: 216 ! Ii protein. Identification of mutant Ii	1: \\\\x(\frac{1}{2})(\frac{1}{
NEGVETLGGSVAGLGAMMIA-AREDEE 1 1 EAR PAGE 1 1 1 1 1 200 1 1 1 1 1 200 1 1 1 1 1 1	
x(28)(I)x(T)(L)xx(V)(T)(R)xx(V)xxx(T)x(28) 1: VPSSNSASQSPOANTDFCSPRLSYLSKAILTLCEVTRKPPEVVTITRWKAPVVWEGTYNRAVL DEFENDENTENDEN	ps3748 ck. 2863 len: 355 ! Seven transmembrane receptor (V28). DNA end
R025U/ CK: 23U3 IEH: 3E3 - GHERCOUJE TOTAL POR A, P)XX(L,I,V,M,A,F,Y,W	ALQTEIANLLKEKEKLEFILAAHRPACKIDDLGFPEEMSVASLDLTGGLEVATP RSREAFTLPLLMDPEFKPSVEFVKSISSMELKTEPFDDFLFPASSRPSGSETARSVPDMDLSGSFYAADWEPLHSGSLGMGPMATELE
	1 <(x){1,200}(L,1,V,M,A,P)X(P,T,S)(L,1,V,M,A,P)XX(L,1,V,M,A,E,T,W)(V,X,99)(A)X(S)(A)X(Y)(S)(B)X(V)XXX(M)X(265) x(99)(A)X(S)(A)X(Y)(S)(B)X(V)XXX(M)X(265)
1 <pre><(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,N,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y</pre>	R53646 ck: 4677 len: 380 ! c-fos gene product. New antitumour vaccines
R63444 ck: 7811 len: 120 ! Amino terminal sequence of the rubisco a	1: x(5)(L)x(S)(A)xx(A)(T)(R)xx(F)xxx(S)x(35) MASSMLSSAAVATRTNPAQASMVAPFTGLKSAAFPVSRKQNLDITSIASNGGRVQC
1 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{4}(Y)x(T)(Y)xx(A)(T)(R)xx(L)xxx(G)x(100) MATAVSTVGAATRAPLNLNGSSAGASVPTSGFLGSSLKKHTNVRFPSSSRTTSMTVKAAEN	R53251 ck: 2035 len: 56 ! Consensus sequence of signal peptide of sma
R63443 ck: 8722 len: 120 ! Amino terminal sequence of the rubisco a	1: RYSAPPGDPKKFILLELIISSSNYSYALISKSNYSYALISKSSNYSYALISKSSNYSYALISKSNYSYA
GVPEBLIKKAPSAGLMEGQTDESEMGTTYEMIDKYLKGEEIPERDKKLLEKHHEKS HHKRQLAIAPPKF	1 <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C)
1 <(x)[1,200](L.I.V.M.A.P)X(P.T.S)(L.I.V.M.A.P)X(L.I.Y.M.A.F.X.W x(158)(L)x(P)(L)x(F)(T)(K)xx(V)xx(G)xf(Z) 1: MQEKIDKIVQVISAGINGAVVGISGGIDSAVVAHLIKRAFPDDSLGLIMPCKSNPKDM	R50082 ck: 108 len: 215 ! NANBH virus antigenic fragment #14. Nucleic
R58587 ck: 9708 len: 246 ! Nicotineamide adenine dinucleotide synth	1; LERYHNKYYCTISKSASILAKKYTEDEMQYLDAHIUSVLAH
1 (X)[1,20](L,1,V,E,F,A(F,E,E,F,A(F,E,E,E)) (X) (Y)(X)(X)(X)(X)(X)(X)(X)(X)(X)(X)(X)(X)(X)	xx(A)(C)(R)xx(P)xxx(A)x(284)
R58636 CK: 3320 Tell: IV M B DX(P.T.S)(L,I,V,M,B,P)XX(L,I,V,M,A,F,Y,W	61 ck. 3747 len. 352
מה מה משנים	_

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<(x){1,200}(L,I,V,M,A,P)X(P,T
T.S)(L.I.V,M,A,P)XX(L,I,V,M,A,F,Y,IXX(V)XXX(G)X(225) XX(V)XXX(G)X(225) IMPETISROCEVLASEPLGIPIVVRSADLYRESSSI REIPAP GDVTWMERLLWGVFGRTSTR
1: PVPSLATISLENSLWSALSKO KNIQLLLNYKQNVA R84103
72715 ck: 516 len: <(X)[1,200](L,1) <(X)[1,200](L)X(T) XX[170](L)X(T) MXENNILSVADLARE CEPOATLIGHTUSEPHLUIGT TPVADIAAQALCPQGINAMED
R70100 ck: 4652 len: 274 ! Lettuce infectious yellows virus open read (X)[1,200](L,I,V,M,A,P)X(P,I,V,M,A,P)X(L,I,V,M,A,F,Y,M)(x(3)(M)x(P)(L)xx(L)(T)(K)xx(Y)xxx(A)x(255) MIMMSEPLYALIRGOVUDTAXYLAVETQHCAITTVACRILELSVGFMTIVKLCGFKMDTSSFIASI KVVTDLYLDLIGVLYNUTGYNIDLVEKEIFDKLEFLVYDGEEGFKSPQVEYNDICT VYNLKPIIKYNRWHTDGSIVIECGDVIGKGINKTKKKICNK
R71324 ck: 7564 len: 329 ! Acetyl-CoA-reductase. Transgenic plant mate <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x(195](L)X(S)(L)X(Y)(T)(K)XX(I)XXX(A)X(IIS) VASMISSAVATSGOSAVATGEPVKKVNTDITSITSNGGRVKCMQVWPPIGK VVFRKMTRADWDAVIDTNLTSLFNVTKOVIDGMADRGWGRVNISSVNGQXQOFGQ TNYSTAKACLHGFTMALAQEVATKGVTVNTVSPGYIATDMYKAIRQDVLDKIVATIPVKRLGLPEEIASICAWLSSEESGFSTGADFS TNYSTAKACLHGFTMALAQEVATKGVTVNTVSPGYIATDMYKAIRQDVLDKIVATIPVKRLGLPEEIASICAWLSSEESGFSTGADFS
R67596 ck: 5465 len: 331 ! A. aculeatus pectin methylesterase. Enzyme <(X) [1, 200] (L, I, V, M, A, P)X(P, T, S) (L, I, V, M, A, P)XX(L, I, V, M, A, F, Y, W) (C x {11] {A}X(S) {A}XX(A)XX(A)X{304} i
R66682 ck: 5012 len: 25 ! Peptide MONOSER. Immunogenic complex contg. (X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x(2)(I)x(P)(A)xx(Y)(S)(R)xx(Y)xxx(A)x(7) 1: CNIAPASIVSRNIVYTSAQPNQDIA
1 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W) x(I)x(P)(A)xx(V)(S)(R)xx(V)xxx(A)x{8} 1: NISPASIVSRNIVYTRAQPNQSIAC

R89897

ck: 206

len: 437

! Rat kynurenine aminotransferase (KAT) cl

1 <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W
x(161)(L)X(P)(A)XX(A)(P)XXX(P)XXX(V)X(246)
1: MYKRIQARKIOSIOQHLWEFGKLIKEYDVVNLGGGFDFSPDFATQAFQQATSGNFMLNQY
LVLNTPNNPLGKVFSRMELELVANLCQQHDVVCISDEVYQMLVTDGHQFYSIASLP
GMWDRTLTIGSAGKSFSATGWKVGWVMGPDNIMKHLRTVHQNSIFHCPTQAQAAVAQCFEREQQHFGQPSSYFLQLPQAMELNRD

R89896 ck: 6122 len: 423 ! Rat kynurenine aminotransferase (KAT) cl

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1 <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W
x(156)(P)X(T)(V)XX(L)(S)(H)XX(L)XXX(A)X(182)
1: YCCRRREWRINKRESTSTATITEPSELLLDRLHPNMYGRMPLLLNPKLLSLEYPRNNIEYV
GPPPLSCAEQLCIARQVAAGMAYLSERKFVHRDLATRNCLYGETMYVKIADFGLLSR
NIYSADYYKADGNDAIPIRMYPPESIFYNRYTTESDVMAYGVVLWEIFSYGLQPYYGMAHEEVIYYVRDGNILACPENCPLELYN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ::
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                         R94990
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         R91274
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R81333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R81330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R81327
                                                                                                                                                                                                                                                                                                                                                                                                                                               ck: 1456 len: 62
                ck: 5928 len: 282
                                                                                                                                                                                                                                      ck: 5431 len: 354 ! Nsk2 receptor intracellular domain (aa5]
                                                                                                                                                                                                                                                                                                                       <(x)\{1,200\}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,X,43)\}(P)X(P)(I)XX(A)(S)(K)XX(P)XXX(M)X\{3\}\\LQHHHQHIILQQQQHHHPPPPAAVLVAAPPADKEQFAAAAAAAPIPIAAASKETPLIDMMS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ck: 824 len: 225 ! Tyrosine phosphatase MPTP-delta insertio
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          <(x)\{1,200\}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,X,G),X,P), \\ x\{69\}(I)x(P)(A)xx(Y)(T)(K)xx(P)xxx(G)x\{57\} \\ MDWTWRVFCLLAVAPGAHSQVQLVQSGAEVKKBGASVKVSCKASGFNIKDTYIHWVRQAPGQR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ck: 9019 len: 142 ! Human VLA-4 reshaped antibody 21.6 light
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W
x{50}(I)X(P)(A)xx(Y)(T)(K)xx(P)xxx(G)x{57}
EVQLQQSGAELVKPGASVKLSCTASGFNIKDTYIHCVKQRPEQGLEWIGRIDPANGYTKYD</pre>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ck: 8013 len: 123 ! Mouse anti-VLA-4 antibody 21.6 heavy cha
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W
X{69}(I)X(P)(A)XX(Y)(T)(K)XX(P)XXX(G)X(55)
MKCSWVMFELMAVVTGVNSEVOLQQSGAELVKPGASVKLSCTASGFNIKDTYIHCVKQRPEQG</pre>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ck: 2629 len: 140
! Nsk2 receptor tyrosine-kinase region (aa
                                                                                                                                                                                                                                                                                                                                                                                                                            l Eimeria gametocyte antigen. Protection o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ! Mouse VLA-4 antibody 21.6 light heavy va
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٢.	OSFVVSGHDYPRKLOACVAAGLDMYYSEVVSRRINDPSRDFPGDV VEVRGKSVSWEGLEQFVSSATYAGFRRVALMVDAASHYSLMSADDLTSALERKYECTVKVNESVSSFLRDVFVWSPH	AIVEASLIAETQ QVILDGDPLLTV
EVIEIGLALAKAIÕI. QEMKAAEQKSKDLEGIMDIVNI	1 <\(x)\[1,200\](L,V,M,A,P)X\(P,T,S)\(L,L,V,M,A,F),X\(L,L,V,M,A,F,X,M)\(Y,X,V,M,A,F,X,M)\(Y,X,V,Y,X,X,Y,Y,X,X,Y,X,X,Y,Y,X,X,Y,Y,X,X,Y,X,X,Y,X,X,Y,X,X,Y,X,X,Y,X,X,Y,X	ч
д.	W00624 ck: 8391 len: 358 ! SacI endonuclease. DNA encoding SacI restri	
DNTDLAVVEKALTYEKFKAWTN W01743		H
1: KYTRGMLFKTNEEKKTVOMMFK	R99576 ck: 2766 len: 83 ! Wasp venom BrhTX-1 subunit (b) partial sequ	
1 R99253	MAASPMQVFSRRRLLLEVWRSSPDWQQDATVTEHVHRIRRKIEEDPTKPTILQTVR GAGYRFDGERA	MAASPMQVFSRRF GAGYRFDGERA
AAPSTTSALTYGPAAAAADAAA	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q) x(179)(A)x(P)(M)xx(F)(S)(R)xx(L)xx(L)xx(P)x(40) x(179)(A)x(P)(M)xx(F)(S)(R)xx(L)xx(L)xx(L)xx(L)xx(L)xx(L)xx(L)x</pre>	н
1:	R99501 ck: 2030 len: 244 ! Nitrilase regulatory factor 244-amino acid	
W00725	<pre><(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,E,I,W)(Y) x{2}(I)x(P)(A)xx(V)(S)(R)xx(V)xxx(A)x{7} 1: CNIAPASIVSRNIVYTRAQPNQDIA</pre>	ם
RRFKQQKYLSAPEREHLASMIHI GGGGGGGGTGCPQQQQAQQQSPF	R99695 ck: 4995 len: 25 ! VP6 binding peptide B. Encapsulated bioacti	
,, ,	MPYGMALTVIGFILPFAALLACYCLLACRLCRODGPAEPVAOER AAFAISFLPFHITKTAYLAVGSTPGVPCTVLEAFAAAYKGTRPFASANSVLDPILFYFTQKKFRRRHELLQKLTA	YDLSPPALATHYM RGKAARMAVVVAA
R83015 c	1 <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,M)(V,X,R)) x{179}(L)x(P)(P)xx(A))(F)xx(A)X(133) x(F)x(B)X(A)X(B)XX(A)X(133) 1. wrwnwgroadchpTrgovyrBNFKOLLLEPTVSAVLAAGLFUNICVITQICTSRRALTRTAVYTI	щ
VGGLETFDSLTDLVEHFKKTGIE QESEDTAKAGFWEEFESLQKQEV	R91225 ck: 7811 len: 328 Human placenta G-protein coupled receptor p	
μ·	DYKDDDDKLHHHHHKLAPTSSTIKALULUGHHUHHKLAPTSSTIKALULUGHHUHHHKLAPTSSTIKALULUGHHUHHHKLAPTSSTIKALULUGHHUHHHKLAPTSSTIKALULUGHHUHHHKLAPTSSTIKALULUGHHUHHHKLAPTSSTIKALULUGHHUHHHKLAPTSSTIKALULUGHHUHHHKLAPTSSTIKALULUGHHUHHHKLAPTSSTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHHKLASTIKALULUGHHUHKASTIKALULUGHHUKASTIKALULUGHHUHKASTIKALULUGHHUHKASTIKALULUGHHUHKASTIKALULUGHHUHKASTIKALULUGHHUHKASTIKALULUGHHUHKASTIKALULUGHHUHKASTIKALULUGHHUHKASTIKALULUGHHUHKASTIKALULUGHHUHKASTIKALULUGHHUHKASTIKALULUGHHUKASTIKALULUGHTIKALULUGHTIKALULUGHTIKALULUGHTIKALULUGHTIKALULUGHTIKALULUGHTIKALULUGHTIKALULUGHTIKATIKALULUGHTIKATIKALULUGHTIKATIKATIKATIKATIKATIKATIKATIKATIKATIKA	PRGWEQLEQCGYPY EQARLALTLAAAE!
R99316 c	<(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q) ×(157)(A)X(T)(A)XX(A)(C)(H)XX(L)XX(T)X(248) ×(157)(A)X(T)(A)XX(A)(C)(H)XX(L)XX(T)X(248)	H
EVIEIGLALAKAIÕTLGEATKSA QEMKAAEQKSKDLEGTMDTVNTV	R95055 ck: 7433 len: 421 ! IL-2-DETA-DGAL4 multidomain protein. Nuclei	
1.	NDEAGAANADEKLLSSIEQAGDIGEKKKIKGSKEPKGAKGIKNNWEGRYSBKTKR SPLTRAHLTEVESRLERLEQLFLLIFPREDLDMILKMDSLQDIKALLFGLFVQDNVNKDAVTDRLASVETDMPLTLRQHRISATSSSE	NDEAGAANADEKLI SPLTRAHLTEVESF
R94586 cl	$<(x)[1,200](L,I,V,M,\lambda,P)X(P,T,S)(L,I,V,M,\lambda,P)X(L,I,V,M,\lambda,F,Y,W)(G,X,P)](A)xx(B)(G)(H)xx(L)xx(T)x[248]$ $=(x,P)[A,X,C](A)xx(A)(C)(H)xx(L)xx(T)x[248]$ $=(x,P)[A,X,C](A)xx(A)(C)(H)xx(L)xx(T)X[248]$	μ.
EVIEIGLALAKAIQTLGEATKSAI QEMKAAEQKSKDLEGTMDTVNTVI	R95054 ck: 9286 len: 342 ! TGF-a-DETA-DGAL4 multidomain protein. Nucle	
1	<pre><(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q x(20)(L)x(S)(L)xx(I)(S)(H)xx(A)xxx(T)x{124} xAAYISSKERKSMKKYSLVTLLSLECISHAKAQTATLTDEYYKKALENTQKLDVAKSQTAESIYES</pre>	1
DQ4580 Ck	R77433 ck: 2940 len: 160 ! Antigen of Rochalimaea henselae. Nucleic ad	
1 <(EVIETGLALAKATOTIGEATKSAL QEMKAAEQKSKDLEGTMDTVNTVN	2(X)[1,200](L,1,V,M,A,P)(X)(X,Y,E),M,X,E)(X,Y,Y,E)(X,Y,E)(1 LKPSPAPKGKLGAS ANLCQQHDVVCISD
R94579 ck	R89898 ck: 7276 len: 457 ! Rat kynurenine aminotransferase (KAT) clone	

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<(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,A,F,Y,W,A,F,Y,M,A,F,Y,M,A,F)X(L,I)XXX(A)(T)(K)XX(L)XXX(A)X(T)(K)XX(L)XXX(A)X(T)(K)XX(L)XXX(A)X(T)(K)XX(L)XXX(A)X(T)(K)XX(L)XXX(A)X(T)(K)XX(L)XXX(A)X(T)(K)XX(L)XXX(A)X(L)XXX(A)XX(L)XXX(A)XX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(L)XXX(
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           k: 2400 len: 488
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ! Chlamydia pneumoniae polypeptide antigen
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ck: 3773 len: 271 ! C. pneumoniae polypeptide antigen (polyp

<(X){1,200}{L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W
x{189}(I)x(T)(L)xx(A)(T)(K)xx(L)xxx(A)x(66)
MSISSSGSDNQKNIMSQYLTSTPQGVPQQDKLSGNETKQIQQTRQGKNTEMESDATIAGASG
%ALSWYASTOADQTNKLGLEKQAIKIDKEREEY
VMIAKGFELPWGPLIN</pre>

<(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W
x{189}(I)x(T)(L)xx(A)(T)(K)xx(L)xxx(A)x{54}
wsisssgddnoknimsgvlttstoggvdodklignetkqiootrogkntemesdatiagasg
salsnyastqaqadqtnklglekqaikidkereby
rvmia</pre> ck: 9728 len: 259 ! C. pneumoniae polypeptide antigen (polyp

<(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W
x{157}(P)x(S)(P)xx(V)(T)(H)xx(V)xxx(G)x(285)
MYRWTHADLSGLDAETLLKGROVHGSELARPSRKNGDFSLSVRVGDQVTHIRIQNSGDFYDL
IEBASGAFVYLROPYYATRVNADLIENTVLELNKK
IEBASGAFVYLROPYYATRVNADLIENTVLELNKK
EVKNLHORLEGORPENKGKNRYKNILPFDHSRVILOGRDSNIFGSDYINANVIKNQLLGPDENA</pre> ck: 4323 len: 458 ! Human SH-PTP1 variant derived from eryth

ck: 5439 len: 371 ! Human thyroid transcription factor-1. 01

<(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W
x{158}(A)x(S)(P)xx(I)(T)(H)xx(P)xxx(P)x{3}}
MSPLIRRILLAALLQLAPAQAPVSQPDAPGHQRKVVSWIDVYTRATCQPREVVVPLTVELMGT
ASSVAKGGA</pre> ck: 1679 len: 207 ! Vascular endothelial growth factor-like

ck: 4145 len: 374 ! Cytoplasmic antiproteinase-2 protein. Hu

ck: 9728 len: 259 ! C. pneumoniae 53 kDa antig**en**. Detection

<(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,X,189)(I)x(I)x(I)x(A)x(A)x(A)x(54)
MSISSSGPDNQKNIMSQVLTSTPQGVPQQDKLSGNETKQIQQTRQGKNTEMESDATIAGASG
GSALSNYASTQAQADQTNKLGLEKQAIKIDKEREEY</pre>

ц	1 NLGEISPQ ESVDNKLA	н		₩	L	1 MAASPMQVFS GAGYRFDGEI	1 VFICHYFP	1 1YELGNPC	1 AAPSTTSA
W20259	W20776 1 NLGBISPOVVKRVSTVLENKLE ESVDNKLAGAIKEMMFTFEDIA	w20886 1:	W20571	W06935	W06943	W13272 1: FSRRLLLEVWRS	W11479 1: TCHYFPGGNMYKDLIYELG	W11481 1: ELGNPCKHNKDCRTKRCSA	W04831 1: LTPGPAAAAADAJ
ck: 5141 len: 108 ! H. pylori secreted or periplasmic protein 2 <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(d	<pre>ck: 6185 len: 276</pre>	<pre>ck: 2872 len: 154 ! H. pylori secreted or periplasmic protein, <(X)(1,200)(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x(54)(L)x(S)(V)xx(V)(S)(R)xx(L)xxx(I)x(84) NNAKNKKGVKSMKTNFYKIKLLFAWCLIIGMFNAPLNADQNTDIKDISPEDMALNSVGLVSRDQI NNAKNKKKGVKSMKTNFYKIKLLFAWCLIIGMFNAPLNADQNTDIKDISPEDMALNSVGLVSRDQI</pre>	<pre>ck: 6963 len: 114 ! H. pylori secreted or periplasmic protein 8 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x{80}(L)x(S)(L)xx(I)(S)(K)xx(L)xxx(A)x{18} mkrpisklkQnflQFkHsfnkHlDKYSLYYRLfNISSIVIGFLIALFSYGAGVILVYPILFLFA</pre>	<pre>ck: 8017 len: 136</pre>	ck: 9316 len: 140 ! CagI locus product 14. Helicobacter pylori <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x{40}(L)x(S)(Y)xx(Y)(S)(R)xx(L)xxx(I)x{84} KRIFIKLNYYLLGVLSLACLQSVNADQNTDIKDISPEDMALNSVGLVSRDQLKIEIPKETLEQKVA	ck: 2018 len: 244 ! Rhodococcus erythropolis SK92-Bl regulatory <(X)[1,200](L,I,V,M,A,P)X(P,T.S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x(179](A)x(P)(M)xx(F)(S)(R)xx(L)xxx(V)x(49) MAGADVHAGGGTNRRARILVVDDEKHYTHVWQLESENFDVVAAADGDAALRQVTESAPDLMVLD SSPDWQQDATVTEHVHRIRRKIEEDPTKPTILQTVR	CK: 9389 len: 221 ! D. immitis venom allergen antigen 5-like pr <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(G x{173}(A)X(T)(L)XX(T)(C)(H)XX(P)XXX(M)X{32} MILLVIFPAIIVVASYECEGGKLTPTERKNIVTQINKYRSRLIRGKLKNKDGYLMPKGKNMLRMF GNPCKHNKDCRTKRCSAKSGLCKK	ck: 5195 len: 205 ! D. immitis mature venom allergen antigen 5- <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x{157}(A)X(T)(L)XX(I)(C)(H)XX(P)XXX(M)X(32} YECEGGKLIPTERKNIVTQINKYRSRLIRGKLKNKDGYLMPKGKNMLRMRWDCKLEKSAQNWANMC AKSGLCKK	W04831 ck: 1679 len: 207 ! Vascular endothelial growth factor-B186. Va <(X){1,200}{L,I,V,M,A,P}X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x{158}(A)x(S)(P)xx(I)(T)(H)xx(P)xxx(P)x(33} MSPLIRRLILAALLQLAPAQAPVSQPDAPGHQRKVVSWIDVYTRATCQPREVVVPLTVELMGTVAK AAPSTTSALTPGPAAAAADAAASSVAKGGA

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VPSL/				1 NEVPSPAJ QIKDVNSI	VPSL		DLSPI								
1 QVPSLATISLENSLWSALSKOI TKNIQLLLNYKQNVA	W25141	W13647	1	W21709 1: NLSLENEAWSLLSKO	sko	W21704	l: PALATHYMPYGMALTVI RMAVVVAAVFGISFLPF	W09433	H ·	W20139	.	W20696	!	W20324	1:
<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,X,S)X(195) x{37}(L)X(S)(A)XX(A)(S)(R)XX(L)XXX(S)X(195) DVRFSLGGSSTSYSKETGDLRKALDSGNGTVNLTILLSSASGASRYTLMTLSNYDGKAITVA IQLAQINNGTFKTPVVITDDKGQRVEITNVTSKVV</pre>	inhibite	<pre>ck: 5362 len: 30</pre>	(X) (3) THE 'A' FIGURE HEROGER BY CLOIM GETTE GETTE (X) (1,20) (L,I,V,M,A,P)X(P,T,S) (L,I,V,M,A,F,Y,W (X) (B) (Y)x(P)(F)X(A) (S) (R)xx(P)xxx(P)x(S) (R) (X) (X) (X) (X) (X) (X) (X) (X) (X) (X	: 7681 len: 250 ! Luffin-B. Inactive precursor of maize x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,YX;08}(L)xx(S)(A)xx(A)(S)(R)xx(L)xxx(S)x{196} VSFSLSGADSKYSKFITALRKALPSKEKVSNIPLLLPSASGASRYILMQLSNYDAKAL LAQTNNGAFRTPVVIIDNKGQRVEITNLASKV	1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V)7)(L)X(S)(A)XX(D)(S)(R)XX(L)XXX(S)X(195) LSGSSSTSYSKETGDLRALDENGTVVNLTILLSSASGASRYTLMTLS NNGTFKTPVVITDDKGQRVEITNVTSKVV	ck: 7673 len: 248 ! Luffin-A. Inactive precursor of maize ri	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(179)(L)x(P)(P)xx(A)(T)(H)xx(P)xxx(A)x(133} newdngtdoalglepttcyyrenergullepuvcavlapalplnicvitqictsrraltriav LSPPALATHYMPYGMALTVIGFLLPFAALLACYCLLACRLCRODGPAEPVAQER KAARMAVVVAAVFGISFLPFHITKTAYLAVRSTPGVPCTVLEAFAAAYKGTRPFASANSVLDPILFYFTQKKFRRPHELLQK</pre>	ck: 8653 len: 328 ! Human placenta purinergic P-2u receptor,	<pre><(X){1,200}{L,I,V,M,A,P}X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{33}{L)X(T)(I)xx(L)(S)(H)xx(V)xxx(P)x{4} MLDFDLVLFGATGDLAMRKLEVSLYEIYISFMVLKTILGLSHRGVRSYPMKSF</pre>	ck: 1367 len: 53 ! H. pylori cytoplasmic protein, 14257751.	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{87}(L)x(S)(L)xx(I)(S)(K)xx(L)xxx(A)x{18} rssekikmkrpisklkqnelqekhsenkhldkyslyyrlenissivigelialesygagvi</pre>	ck: 7893 len: 121 ! H. pylori secreted or periplasmic protei	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{2}(A)x(S)(L)xx(V)(S)(K)xx(L)xxx(L)x{84} MMAHSLILVSKTSLSNLLIFVVQPDGKLSMTDAAIDPNMTNSGLRWYRVNEIAEKFKLIKD</pre>	ck: 6372 len: 102 ! H. pylori cell envelope transporter prot	x{42}(L)x(S)(V)xx(V)(S)(R)xx(L)xxx(L)x(50) MKTNFYKIKLLFAWCLIIGMFNAPLNADQNTDIKDISPEDMALNSVGLVSRDQLKIEIPKE

μ	н		н		1 YSKALGVI IGMQQMGI		н		ч		1		μ		ч		1: NEVPSPAALSLENEAWS QIKDVNSKLLLNKQNIA	
W2449:	i	w24492	!	W24491	1; NYENLIVCQPDNG QARLMSQALRKM	W34668	1:	W17971	1:	W26413	1:	W22428	1:	W22413	1:	W22410	LLSKC	W25146
<pre>3 ck: 3320 len: 124 ! Novel amylase inhibitor protein 3 subunit i <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,E,Y,W)(C) x{91}(A)x(S)(I)xx(V)(C)(R)xx(I)xxx(A)x{17}</pre>	<pre><(X) [1,200] (L, I, V, M, A, P)X(P, T x(91] (A)x(S) (I)xx(V) (C)(K)X SGPWMCYPGYAFKVPALPGCRPVLKLC</pre>	0	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x(91)(A)x(S)(I)xx(V)(C)(K)xx(I)xxx(A)x{15} sgpwmcypgyafkvpalpgcrpvlklqcngsqvpravlrdccqqladisewcrcgalysmldsm</pre>	ck: 1423 len: 122 ! Novel amylase inhibitor protein subun	1	ck: 9097 len: 438 ! Arabidopsis thaliana Rec-A protein. DNA end	<pre><(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(\ x(23)(L)X(T)(L)XX(L)(C)(R)XX(V)XXX(L)X(8) NSARARASAAPRPGAMLRSCAARLRTLGALCRPPYGRRLPEATRDPS</pre>	inal binding pro	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q,X(83)(A)x(S)(A)xx(F)(T)(H)xx(Y)xxx(L)x(22) x(83)(A)x(S)(A)xx(F)(T)(H)xx(Y)xxx(L)x(22) MIIDVITSLDGGCNKCLNIKHAKSQCKPSSLLISSLENVNPGINPRFFIQNIAANEDEKNIPSI MIIDVITSLDGGCNKCLNIKHAKSQCKPSSLLISSLENVNPGINPRFFIQNIAANEDEKNIPSI</pre>	coded protei	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q x{69}(I)X(P)(A)xx(Y)(T)(K)xx(P)xxx(G)X(57) xxmmrvfcllavapgahsqvqlvqsgabvkkpgasvkvsckasgfnikdtythwvrqapgqrlew</pre>	antibody 21	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q x{50}(I)X(P)(A)xx(Y)(T)(K)xx(P)xxx(G)x(57) qvqLvqSgAEvkkpgASvkvSCKASGFNIKDTYIHWVRQAPGQRLEWMGRIDPANGYTKYDPKF</pre>	ck: 9557 len: 123 Humanised alpha-4 integrin antibody 21.6 VH	<pre><(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(\ x{69}(I)x(P)(A)xx(Y)(T)(K)xx(P)xxx(G)x{55} wkcswymfflmavytgvnsevqlqosgaelvkpgasvklsctasgenikdtythcvkqrpeoglew</pre>	. 4	<pre><(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)X(L,I,V,M,A,F,Y,W)(\) x(38}(L)x(S)(A)xX(A)(S)(R)xx(L)xxx(S)x1196} ANVSSSLSGADSRSYSKFITALERALESSEKYSNIPLLESASGASEXILMQLSNYDAKAITMAIC BLOCK ANVSSSLOPE AND AND AND AND AND AND AND AND AND AND</pre>	proteir

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1 <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,X,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,W,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,Y,X)(L,I,V,M,A,F,X,X)(L,I,V,M,A,F,X,X)(L,I,V,M,A,F,X,X)(L,I,V,M,A,F,X,X)(L,I,V,M,A,F,X,X)(L,I,V,M,A,F,X,X)(L,I,V,M,A,F,X,X)(L,I,V,M,A,F,X,X)(L,I,V,M,A,F,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X,X)(L,I,X,X,X,X)(L,I,X,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I,X,X,X)(L,I
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                                                                                                                                                                                                                                                                                                                     W47606
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SGPWMCYPGYAFKVPALPGCRPVLKLQCNGSQVPEAVLRDCCQQLADISEWCRCGALYSML</pre>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,X)(1,X)x(1,X)x(1,X)x(L,I,V,M,A,F,Y,W,X)(1,X)x(1,X)x(1,X)x(L,I,V,M,A,F,Y,W,X)</pre>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ck: 2685 len: 124 ! Wheat amylase inhibitor 0.26 Ala subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ck: 3320 len: 124 ! Wheat amylase inhibitor 0.19 Al subunit.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ck: 1423 len: 122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W
x(91)(A)x(S)(I)xx(V)(C)(R)xx(I)xxx(A)x(IT)
SGPWMCYPGQAFQVPALPACRPLLRLQCNGSQVPEAVLRDCCQQLAHISEWCRCGALYSML</pre>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        \begin{array}{l} <\langle \mathbf{x} \rangle \{1,200\} (L,L,V,M,A,P) \mathbf{X} \{P,T,S) \{L,L,V,M,A,P) \mathbf{X} \{L,L,V,M,A,F,Y,M,Y,L,L,S\} \} \\ \times \{1.15\} \{P\} \times (\mathbf{T}) \{A\} \times (\mathbf{Y}) \{C\} \{R\} \times (P) \times (P) \times (P) \times (P) \\ + (LL,V,M,A,P) \times (P) \times (P) \times (P) \times (P) \times (P) \\ + (LL,V,M,A,P) \times (P) \times (P) \times (P) \times (P) \times (P) \\ \times (P) \times (P) \times (P) \times (P) \times (P) \times (P) \\ + (P) \times (P) \times (P) \times (P) \times (P) \times (P) \times (P) \\ + (P) \times (P) \times
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ck: 9381 len: 176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W
x(70)(I)x(P)(L)xx(F)(S)(H)xx(A)xxx(I)x(21)
STENLEDANKMLLEISQLSTPFXTSAQRVAAYFSEAISARLVSSCLGIYATLPHTHQSHKV</pre>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ck: 1840 len: 107
<(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,X)
x{83}(I)x(S)(I)xx(I)(T)(K)xx(A)xxx(M)x{95}
mSRRNPCKFEIRGHCLNGKRCHFSHNYFEWPPHALLVRQNFMLNRILKSMDKSIDTLSEISGA</pre>
                                                                                                                                                                                                                                                                                                                                        ck: 9828 len: 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SGPWMCYPGQAFQVPALPACRPLLRLQCNGSQVPEAVLRDCCQQLAHISEWCRCGALYSML
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ! Residues 217-392 of human type B EBNA2 (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ! Wheat amylase inhibitor 0.26 Alb subunit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ! Soybean SCARECROW SRPgl protein. DNA enc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ! Streptomyces ketoreductase (frenolicin 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ! Hepatitis C virus antigen. Protein contg
                                                                                                                                                                                                                                                                                                                                        ! HRSV protein 22K. Production of human re
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein luf
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 \vdash

<(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x{68}(L)x(T)(A)xx(A)(S)(R)xx(L)xxx(P)x{255}	<pre><(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C)</pre>	w59024 ck: 9911 len: 71 ! Enterocin-900. Bacterial growth inhibiting	1 <(X){1,200}(L,I,V,M,A,P)X(P,I,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C 1 x{170}(P)X(T)(A)xx(V)(T)(X)xx(A)xxx(M)x{131} x{170}(P)X(T)(A)xx(V)(T)(X)xx(A)xxx(M)x{131} xxx(M)x{131} xxx(M)x{131} xxx(M)x{131} xxx(M)x{131} xxx(M)x{131} xxx(M)x{131} xxx(M)x{131} xxx(M)xx{131} xxx{131} xxx{131	W37816 ck: 3813 len: 317 ! Human secreted apoptosis-related protein hs	W28052 ck: 2273 len: 269 ! Amino acid sequence of phospho-beta-glucosi <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x(4)](I)x(P)(V)xx(L)(S)(H)xx(M)xxx(L)x(210) MGLKCLRTSIAWTRIFPNGDEDVPNEEGLAFYDRIFDELIAQGIEPVVTLSHFEMPLHLAKHYGGF AQIANRLRFFFPDVQVRGYYPSYKKMLARKGYDVGWQEGDDSILQQGTXDYIGFS YYMSTAVKHDVDTTXENNIVNGGFGFICGESAYRNE AVALGS	W51011 ck: 7216 len: 277 ! Human liver carbonyl reductase. DNA encodin <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x{189}(P)x(S)(A)xx(Y)(T)(K)xx(Y)xxx(S)x172} 1:	W55137 ck: 7408 len: 186 ! Open reading frame 2 peptide of bacteriopha (X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x{74}(L)x(P)(M)xx(F)(S)(K)xx(A)xx(A)x{96} 1: MNPEQLINTLGSAVSGVAQGLSFLPGIASGVLGYLGAQKQNATAKQIAREQMAFQERMSNTAYQRAKNLFSRKGR	W55815 ck: 5054 len: 271 i Streptomyces roseofulvus frenolicin gene cl (X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x(125)(L)x(S)(V)xx(V)(T)(R)xx(L)xxx(G)x(130) 1: MTTAABHTRPGEAGTTRGPALVTGATRGIGLAVAEALVARGYPVVVCARDAEAVARTVKELAAGGA TKALGLELARSGITVNAVCPGYVETPAAAGVRHYADLMDVTEEDVLARFEAKIPL GRYTRPDEVAALVDYLVTDAAAAVTAQALNVCGGLGNY LLVVIVFFIL	1 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x{187}(A)X(T)(L)XX(A)(S)(R)XX(I)XXX(S)X[104} 1: MKKVLLITLSLSLSFWLHAERNGFIVIGLNFAERSYIQGQGSIGEKASAENALNQAINNAKNSLFPN YKDMTGRTLDADTLKRASRHIIRKSSGLVIGMETASANNLTPFNQVKSRTI FQLQGKFGVRFSSDEYDIDRYGDENYLGGSSVELGVKVPAFKVNYYSDDYGDKLDYKRVVSVYLNYTYNFKNKH 1 FQLQGKFGVRFSSDEYDIDRYGDENYLGGSSVELGVKVPAFKVNYYSDDYGDKLDYKRVVSVYLNYTYNFKNKH	X{61}(V)x(P)(A)xx(I)5(X)x(A)xxx(I)x{288} HOTPRETISERLSCVQDKILIDHYENDSKOLDSQIQYWQLIRWENAIFFAAREHGIQTLNHQVVPAY SECEKYGNTGTWEVHFGNNVIDCNDSMCSTSDDTVSATQLVKQLCHTPSPYSSTVS VGTAKTYGQTSAATRPGHCGLAEKQHCGPVNPLLGAATPTGNNKRRKLCSGNTTPTIHLKGDRNSLKCLRYRLRKHSDHYRDISSTWH 1 W55590 ck: 428 len: 307 ! H. pylor1 ORF 09cpl0224_429510_c2_46.aa cel	Ω
W44905 ck: 1135 len: 39 ! "polyproline beta-turn helix" polypeptid	1 < <(x) {1,200} (L,I,V,M,A,P)X(P,T,S) (L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x {170} (V)x(P) (V)xx(I) (X) (H)xx(A)xxx(A)x {85} QSTHPCAAFHAASRALSGGPIYVSDSVGQHDFALLRRLALPDGTVLRCEGHALPTRDCLFADP SHERAIKFAPIGLANMLNTAGAVQAFEAKKDASGVTAEVFVKGAGELVAYSATPR LCKVNGDEAEFTYKDGVVTVDVPWSGSSSKLCCVQYVY	W57889 ck: 5290 len: 271 ! Corn raffinose synthetase. New nucleic a	1	W60764 ck: 7422 len: 260 ! Rainbow trout interleukin 1 beta. Rainbo	W60267 ck: 843 len: 387 ! Klebsiella pneumoniae DHAT protein. New <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(143)(A)x(T)(A)xx(V)(T)(R)xx(V)xxx(T)x(228) MSYRMEDYLVPNVNFFGPNAISVVGERCQLLGGKKALLVTDKGLRAIKDGAVDKTLHYLREAG INDPLLMIGKPAALTAATGMDALTHAVERYISKDANPVTDAAAMQAIKHLARNLRQ INDPLLMIGKPAALTAATGMDAFNNANLGYVHAMAHQLGGLYDMPHGVANAVLLPHVARYNLIANPEKFADIAELMGENITGL	W60258 ck: 4443 len: 339 ! klebsiella pneumoniae glycerol-3-phospha <(x)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)X(L,I,V,M,A,F,Y,W x(68)(L)X(T)(A)XX(A)(S)(R)XX(L)XXX(P)X(255)	<pre>w54728 ck: 3126 len: 24 ! Peptide from Ii p80-103. Increasing upta <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(P)x(P)(x)xx(V)(S)(K)xx(M)xxx(L)x{7} 1: LPKPPKPVSKMRMATPLLMQALPM</pre>	W48722 ck: 2863 len: 355 ! Human V28 seven transmembrane receptor. <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(80)(V)x(T)(L)xx(W)(T)(H)xxX(I)xxX(G)x(259) 1: MDQTERSVTERNETVDLABACYIGDIVVECTVELSIFYSVIFAIGLVGNLLVVFALTNSKKPK TQBIWPVLRNVETNFLGFLLLIMSYCYFRIIQTLFSCKNHKKAKAIKLI VFFLFWTPYNVMIFLETLKLYDFFPSCDMRKDLRALSVTEFVAFSHCCLNPLIYAFAGEKFRRYLYHLYGKCLAVLCGR	W55052 ck: 4306 len: 118 ! Sunflower antifungal protein MS59 fragme <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W X{41}(I)X(T)(A)XX(Y)(S)(H)XX(A)XXX(C)X(61) 1: DPSFPITGEVYTPGNSSFPTVLQNYIRNLRFNETTTPKPFLIITAEHVSHIQAAVVCGKQN	. W62641 ck: 7060 len: 137 ! Flea serine protease inhibitor variable <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{88}(A)x(T)(A)xx(A)(T)(R)xx(I)xxx(A)x{3} 1: EEKLQNVDLQNLTQRNYSVEVILDLPKFKIESEINLNDPLKKLGMSDMFVPGKADFKGLLEGS	GADPATFMGMAGLGDLVLTCTDNQSRNRRFGMMLGQGMDVQSAQEKIGQVVEGYRNTKEVRELAHRFGVEMPITEEIYQVLYCGK

1	1
W60267 ck: 843 len: 387 ! Klebsiella pneumoniae DHAT protein. New	
1 <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x{68}(L)x(T)(A)xx(A)(S)(R)xx(L)xxx(P)x{255} mnornamtvicagsyctalatilarrghevvimghdpehiatlerdrcnaafledvefedtl rvysnedfigvolggavknviaigagmsdgigfganartalitrglaemsrlgaal gaddatfmgmaglddlvltctdnosrnrrfgmmlgogmdvosagekigovvegyrntkevrelahrfgvempiteeivovlycgk	1 RVYSNPDF: GADPATFM
W60258 ck: 4443 len: 339 ! Klebsiella pneumoniae glycerol-3-phospha	
<pre><(X){1,200}{L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(P)x(P)(P)xx(V)(S)(K)xx(M)xxx(L)x(7) LPKPPKPVSKMRMATPLLMQALPM</pre>	н
W54728 ck: 3126 len: 24 ! Peptide from Ii p80-103. Increasing upta	
1	1 DYPEVLQE: LLVVIVFF)
W48722 ck: 2863 len: 355 ! Human V28 seven transmembrane receptor.	
<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(41)(I)x(T)(A)xx(V)(S)(H)xx(A)xxx(C)x(61} 1: DPSFPITGEVYTPGNSSFPTVLQNYIRNLRENETTTPKPFLIITAEHVSHIQAAVVCGKON</pre>	н
W55052 ck: 4306 len: 118 ! Sunflower antifungal protein MS59 fragme	
<pre><(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W x(88}(A)X(T)(A)XX(A)(T)(B)XX(I)XXX(A)X(33} 1: EEKLQNVDLQNLTQRMYSVEVILDLPKFKIESEINLNDPLKKLGMSDMFVPGKADFKGLLEGS</pre>	ч
. W62641 ck: 7060 len: 137 ! Flea serine protease inhibitor variable	•

AVALGSNIQARENNAYASLLAGMAR NIGHTANA	MSYRMEDYLVPNYREFERNANGAIGHTARNIKG INDELLMIGKPAALTAATGMDALTHAVEAYISKDANEVITDAAAMQAIRITTARNIKG INDELLMIGKPAALTAATGMDALTHAVEAYISKORTITDAAAMQAIRITTARNIKG	w30679 ck: 843 len: 387 1,3-Propanediol oxidoreductase. Fermentativ $(x)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,M)(S) \\ \times (1,200](L,I,V,M,A,P)X(P,Y,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,M)(S) \\ \times (1,200](L,I,V,M,A,P)X(V,Y,X,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,$	VSVETEVAEVKGEVRENVMTYYVSYAVEELSSELVSSCOSSELVSSCOSSCORGVLLVSMVVLETEAILCTETRNRILE ASLSYMVGMIASEYNTEAVIMAVGITTAVCETVVIESMQTRYDETSCMGVLLVSMVVLETEAILCTETRNRILE	<pre>-(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,F)A(L,I,V,M,A,F)A(L,I,V,M,A,F)A(L,I,V,M,A,F)A(L,I,I,V,M,A,F)A(L,I,I,I,I,I,I,I,I,I,I,I,I,I,I,I,I,I,I,I</pre>	W62612 ck: 8912 len: 369 ! Human glutamate-binding protein	1 <(X){1,200}(b,1,7,0,7,6,7,7,7,7,7,7,7,7,7,7,7,7,7,7,7,7	W44882 ck: 8918 len: 61 ! "polyproline beta-turn helix"	1: x{27}(P)X(T)(M)XX(H)(X)(M)XX(H)(TMFSLTRQVLNVGPRVPIGPNP) 1: EFTDAGKKANWDGPKSWGLRLYRTGTDPTTMFSLTRQVLNVGPRVPIGPNP	W44879 ck: 9655 len: 104 ! "polyproline beta-turn helix" polypeptide ! <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)((1	44910	1	H X	1 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(V,Y,Y,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,Z,	W44908 ck: 1205 len: 79 ! "polyproline beta-turn helix" po	1 (X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Y,Y,S)(E,Y,X,Y,S)XX(L,Y,X,Y,S)XX(L,Y,Y,M,A,F,Y,Y,M)(Y,S)XX(L,Y,X,X,P)X(L,S,X,Y,S)XX(L,Y,X,S)XX(L,Y,X,S)X(L,Y,X	W44907 ck: 3740 len: 127 ! "polyproline beta-turn helix" po	<pre>(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,1,V,M,A,F)^^(V,T,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,</pre>	W44906 ck: 6449 len: 70 "polyproline beta-turn helix" po	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,P)X(A){D}X{A}{P}X{A}{P}X{D}XX(L)XXX(P)X{D}XXX(P)X{D}XX(L,I,V,M,A,P)XX(L,I,XXXX(P)XIA)</pre> <pre>1:</pre> <pre>AAADPITMESLIRQVLNYGPRVPIGPNPVLPDQRLPSSA</pre>	
eic acid encod , 1 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W	w61619 ck: 1162 len: 250 ! Clone HTPEF86 of TM4SF superfamily. New	-(x)(1,200)(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,F)X(P,T,S)(L,I,V,M,A,F)X(F,T,S)(L,I,X,X,V)X(ZS)		1: MVTHSKEPAACMSREDDTSLÆLKTESSKSEYQLVVNAVRKLQESGEYMSAVIGGEANHUHSK LSRPLSSNVATLQHLCRKTVNGHLDSYEKVTQLEGETREETLDQYDAPL	بر ج	IWGLPLPTRLKDYLEEYKFQV) (C 1	W62614	polypeptide H * x{171}(V)x(P)(L)xx(L)(C)(K)xx(L)(C)(K)xx(L)(C)(K)xx(L)(C)(K)xx(L)(C)(K)xx(L)(C)(K)(K)(K)(K)(K)(K)(K)(K)(K)(K)(K)(K)(K)	E	1	IGPNEVLEDC W62624	ypeptide F A,F,Y,W)(C PNRSKRWNPPRSEGLPAGHHQO	(,A,F,Y,W)(\)	, <u>, , , , , , , , , , , , , , , , , , </u>	ATRGGRCNPLV	1: LCRQRIVAAVGRENLARIPLD	W6261/ Cx: July LU: LU:	peptide D GLAGHVLLDLGALTE	A, E, Y,	

*(62) TMETPRE TMETPRE KSECEKYGNTGTWEVHFGNNVIDCNDSMCI SVGTAKTYGQTSAATRPGHCGLAEKQHCG	<pre><(X)[1,200](L,1,V,M,A,P)X(F,T,S)(L,1,V,M,A,P)XX(L,1,V,M,A,F,</pre>	
	W69891 ck: 1805 len: 375 ! Pig growth differentiation factor-8. Transg	
VRIGKFEFEFVD 1: MSASEGRI VRIGKFEFEFVD W68494 ck: 998:	<pre>1</pre>	
W80624 ck: 450	W69889 ck: 1496 len: 376 ! Rat growth differentiation factor-8. Transq	
1 <(X){1,: x(17)g 1: MAGADYH MAASPMQVFSRRRLLLEVWRSSPDWQQDA: GAGYRFDGERA	W79302 ck: 8790 len: 67 ! A Staphylococcus aureus protein. New nuclei <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C	
GHDLAVITPEPGEDGLTPFLEVKVIDTPK W80799 ck: 2010	1 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(d x{53}(P)x(T)(A)xx(Y)(S)(R)xx(I)xxx(T)x{92} nkyddamalygsdkpdtredmilqdltevvkgvdfkvfsbalavkaivvnglepltadnysrkdid	
X(55) 1: MQKLQIS KPMKDGTRYTGIRSLKLDMNPGTGIWQSII	W38716 ck: 3773 len: 161 ! S. pneumoniae aspartyl tRNA synthetase. Nov	
w69887 ck: 930	1 <(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C x{76}(A)x(T)(P)xx(A)(S)(X)xx(I)xxx(I)x{8} 1: vmpxvfwxpfpxifigtalplagtvatgaihftanevipigxmlxnngliainlayonldraffy	
GHDLAVTFPGPGEDGLNPFLEVKVTDTPK	W77666 ck: 5825 len: 100 ! Staphylococcus aureus protein of unknown fu	
W69886 CK: 146 (X){1,: X(65) MOKIQIC: KDMXTGTBYTGTBSIXITMNDGTGTWSTCTWSTCTWSTCTWSTCTWSTCTWSTCTWSTCTW		
}	W70962 ck: 4556 len: 212 ! A STAT function regulatory protein designat	
W69885 ck: 181 1 <(X){1,:	W38580 ck: 5404 len: 174 Streptococcus pneumoniae protein of unknown <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(G x{72}(V)x(T)(V)xx(L)(S)(K)xx(V)xxx(I)x{86} 1: MGFNQPLSWLILHAKGLTESFIKASAQTGADRITYISCNVRNHGAVDIKLYQELGYELKKVQPVDL	
<(X) x MMQH SPGTG1	W64373 ck: 3575 len: 120 ! Mycobacterium tuberculosis antigen RDIF5. N <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x(95](A)x(S)(A)xx(A)(C)(R)xx(P)xxx(P)x(P) 1: TPEKSFVDDLDIDSLSMVEIAVQTEDKYGVKIPDEDLAGLRTVGDVVAXIQKLEEENPEAAQAL	
W30689 ck: 229	KPMRUGTKTIGIKSIKLUMNEGTGIMUSIUVKIY LUMMAKQEBSNILGIEIKALUEN GHDLAVTFPGPGEDGLNPFLEVKVTDTPKRSRRDFGLDCDEHSTESRCCRYPLTVDFEAFGWDWIIAPKRYKANYCSGECEFVFLQKY	
MOKLOIF KPMKDGTRYTGIRSLKLDMNPGIGIWQSII GHDLAVTFPEPGEEGLNPFLEVKVTDTPK	<pre></pre> <pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,X)(L,S)(L,X,Y,M,A,P)XX(L,I,V,M,A,F,Y,X)(DS)(K)XX(I)XXX(L)X{294} MQKLQCYYYLFMLIVAGPDLNENSEQKENDENENGCTWRQNTKSSRIEAIKIQILS DOTGINGSTINGSTYLONGI, KODESNIGITETETIDEN</pre>	
1 <(X)[1,: x(65)	W65460 ck: 1814 len: 375 ! Human growth differentiation factor-8. Tran	
GHDLAVTEPGPGEDGLNPELEVKVTDTPKI W69892 ck: 122	x{23}{P}x(T){P}xx(N)(S)(H)xx(L)xxx(S)x{211} NNSMTSAVPVANSVLVVAPHNGYPVTPGIMSHVPLYPNSQPQVHLVPGNPPSLVSNVNGQPVQKAL VNPGMAISGVLLVFCLLEFGIACASSHFGCQLVCCQSSNVSVIYPNIYAANPVITP EPVTSPPSYSSEIQANK	

KRSRRDFGLDCDEHSTESRCCRYPLTVDFEAFGWDWIIAPKRYKANYCSGECEFVFL len: 375 ! Ovine growth differentiation factor-8. T

,200)(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,I)(L)x(T)(A)xx(I)(S)(K)xx(I)xxx(L)x(294)
ITVYIYLFMLLVAGPVDLNENSEQKENVEKKGLCNACLWRQNNKSSRLEAIKIQILSK
IDVKTVLQNKLKQPESNLGIEIKALDEN
KRSRRDFGLDCDEHSTESRCCRYPLTVDFEAFGWDWIIAPKRYKANYCSGECEFLFL

376 ! Murine growth differentiation factor-8.

.,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W);(L)X(T)(A)xx(I)(S)(K)xx(I)xxx(L)x{294}
)MYVYIYLFMLIAAGPVDLNEGSEREENVEKEGLCNACAWRQNTRYSRIEAIKIQILS
)SIIDVXIYLQNWLKQDESNLGIEIKALDE
:PKRSRRDFGLDCDEHSTESRCCRYPLTVDFEAFGWDWIIAPKRYKANYCSGECEFVF

len: 375 ! Human growth differentiation factor-8. T

., 200)(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(L),I,V,M,A,F,Y,W)(L),I,XX(I),XX,I)(S)(K)XX(I)XXX(L)X{294}
CVYIYLFMLIVAGPVDLNENSEQKENVEKEGLCNACTWRQNTKSSRIEAIKIQILSK
SIDVKTVLQNKIXQPESNICITEIKALDEN
%KRSRRDFGLDCDEHSTESRCCRYPLTVDFEAFGWDWIIAPKRYKANYCSGECEFVFL

63 len: 375 ! Baboon growth differentiation factor-8.

.,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,I)(I)X(I)(A)XX(I)(S)(K)XX(I)XXX(L)X{294}
CVYIYLFMLIVAGPVDLNENSEQKENVEKEGLCNACTWRQNTKSSRIEAIKIQILSK
ILDVKTVLQNWLKQPESNLGIEIKALDEN
KRSRRDFGLDCDEHSTESRCCRYPLTVDFEALGWDWIIAPKRYKANYCSGEGEFVFL

.,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,I)(L)x(T)(A)xx(I)(S)(K)xx(I)xxx(L)x{294}
SVYIYLFMLIAAGPVDLNENSEQKENVEKEGLCNACLMRENTTSSRLEAIKIQILSK
ILDVKTVLQNMLKQPESNLGIEIKALDEN
KRSRRDFGLDCDEHSTESRCCRYPLTVDFEAFGWDWIIAPKRYKANYCSGECEFVFL

len: 375

! Bovine growth differentiation factor-8.

l8 len: 244 ! Rhodococcus nitrile hydratase gene fragm

.,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W))(A)x(P)(M)xx(F)(S)(R)xx(L)xxx(V)x{49} |HAQGGTHRRAILUYUDEKHYPIMYUDESENFDVVAAADGDAALRQVTESAPDLM |ATVTEHVHRIRRKIEEDPTKPTILOTVR

189 1 S. pneumoniae GTP binding protein. Strep

.,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W }(I)x(P)(I)xx(L)(T)(K)xx(L)xxx(L)x{106} RDPYEDYLAINKELESYNLRLMERPQIIVTNKMDMPESQENLEEFKKKLAENYDEFE 82 len: 368 ! E2 papillomavirus protein encoded by pla

.,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,E,Y,W)
[(Y)x(P)(A)xx(I)(S)(K)xx(A)xxx(I)x(290)
[ETLSERLSCYODKIIDHENDSKOIDSIQYWQLIRWENAIFFAAREHGIQTLNHQV
[CSTSDDTYSATQLYKQLQHTPSPYSSTV

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NGHDLAVTFPGPGEDGLNPFLEVKVTDTPKRSRRDFGLDCDEHSTESRCCRYPLTVDFEAFGWDWIIAPKRYKANYCSGECEFVF	Y01440 ck: 9731 len: 64 ! Secreted protein encoded by gene 58 clone H
W97886 ck: 2420 len: 376 ! Murine myostatin. Increasing musc <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M, x{66}(L)X(T)(A)XX(I)(S)(K)XX(I)XXX(L)X(294) 1: MMQKLOMYVYIYLFMLIAAGPVDLNEGSEREENVEKELLNACAMFONTOPYCTER	X{115}(I)X(T)(V)XX(F)(S)(K)XX(I)XXX(S)X{205} MNUKKVLMDISWSUKGGIGFETDEISKLLCDISKEELYRKCASPLAPLGLAVNIFLRKKTDVPELF VISAFAKADIDPSIKLVFTGNPCNDLEKLIIQHGLSERVKFFGFVSEKDLPSLYKG SLGLVFPSLYEGFGLPVVEGMACGIPVLTSLTSSLPEVAGDAALLVDPLSEDAITKGISRLINDSELRKHLIQKGLLRAKRFNWQNVV
GHDLAVTFPEPGEDGLTPFLEVKVTDTPKRSRRDFGLDCDRISMLSLPSNCGF	W88330 ck: 7734 len: 336 ! Mannosyl transferase involved in O antigen (X){1,200}(L;I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XY/I TY, X, T, T, Y, Y, T, Y, T, Y, T, Y, T, Y, T, Y, Y, T, Y, T, Y,
W97885 ck: 1462 len: 286 ! Bovine myostatin (mutant f <(X)[1,200](L,T,V,M,A,P)X(P,T,S)(L,T,V,M,A,P)XX(L x(55](L)X(T)(A)XX(I)(S)(K)XX(I)XXX(L)X;205] MOKLQISVYILLEMLIVAGEVOLMENSEOKENVEKEICHAFTINE	V,M,A,P)X(P,T,S)(L,I,V,)xx(A)(C)(K)xx(P)xxx(S) LSLFSACKDEPKKSSQSHQNNTK
AWY/884 CK: 8371 len: 375 ! Bovine myostatin. Increasing muscle mass (X)[1,20](L], V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,M,G))X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,M,G)XX(L)X(294) KPMKDGTRYTGIRSLKLDMNPGTGICOSIDVKTVLONMLKOPESNLGIEJKALDEN GHDLAVTFPEPGEDGLTPFLEVKVTDTPKKSRRDFGLDCDEHSTESRCCRYPLTVDFEAFGMDMIIAPKRYKANYCSGECEFVFL	New isolat
ин "-	W98862 ck: 4789 len: 304 ! H. pylori GHPO 1670 protein. New isolated H <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) X(L)X(S)(A)XX(Y)(C)(K)XX(A)XXX(L)X{287} **X(L)X(S)(A)XX(Y)(C)(K)XX(A)XXX(L)X{287} **X(L)X(S)(A)XX(Y)(C)(K)XX(A)XXX(L)X(S) **MESALKYCKEKAIDLLVGFVRTYMAQECNILGIYDDAFIITKQENLVGIISLQGLSYSNLMQK **INSKKNKFKKKITFSTKSKKALLIQTIERVKNALKEFKFTLLNSKEVLNFYA EYINGKYIAFNPKLKTSDTILHLMCILRKITLSLNFKIKTPFVRVMGLRLMFAKKFLRSLYLLFYTPKLN
.062 ck: 4555 len: 116 ! H. Pylori ORF 06cp30603_10744075_c <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,P)XX(L,I,V,M,A,P)XX(L,I,V,M,A,P)XX(S)X(S)X{86}	A C
Y11009 ck: 2840 len: 142 ! H. pylori ORF hp6p22217_23564012_f1_5 se 1 <(X)(1,200)(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,X,L)xxx(L)x(B4) 1: MKTNFYKIKLLFAWCLIIGMFNAPLNADQNTDIKDISPEDMALNSVGLVSRDQLKIEIPKETL	<pre>(X)[1,700](L,1,7,M,A) <(X)[1,700](L,1,7,M,A) x[88](A)x(T)(Y)xx(M) MSRPLGFIGYEEGDDEMFVQQMI ACCUMULATION CONTROL ACCUMULATION A</pre>
W97722 ck: 1929 len: 302 ! Staphylococcus aureus mutant P15B9 v1rul - (X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W, M,	9823 len: 221 Se){1,200}{L,I,V,M,A,P}} (160){M,XX(M){K} GIFATTPGIGAFGNKPETYEFVK QMLFDLESAYNAFNKFLHA QMLFDLESAYNAFNKFLHA
1	W89100 ck: 1342 len: 36 ! Polypeptide fragment encoded by gene 206. N <(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C) x{5}(P)x(T)(L)xx(I)(S)(R)xx(L)xxx(P)x{15} 1: FSVHRPETLENISRFLLHSLPKDIPSGISKVKILFT
<pre><(X){1,200}(L,I, x(A)x(S)(V)xx(MAASVGRATRSAAA</pre>	1 <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q) x(112)(A)x(S)(A)xx(I)(S)(H)xx(A)xxx(T)x(223) RPOYCEAMLFALALQTOYCYGIRWLALSKTPSALALANTOTOHCKQLEGIVSAQVQLCRSNLELMHTV TGSQANKLMGLHNSEVGRQALRASLEMKCKCHGYGGHCYIGTSWGLOELQVYAALALNQTQHCKQLEGIVSAQVQLCRSNLELMHTV LKTRYLSATKVVHRPMGTRKHLVPKDLDIRPVKDSELVYLQSSPDFCMKNEKVGSHGTQDRQCNKTSNGSDSCDLMCCGRGYNPYTDH

н	PWFEAAGRLAG PKWNIAVTGN	.	r		ц		GGGGPGGGLLGG AAAAAAAAGQVA	μ		1		н		н		ŀ		RNGNRRLVVTEENI		
Y07041	1: LGPIIDPAF	Y05545	1:	Y11993	1:	ф Оп Э	1: SAHPHPHMH AASAAAAVV		W96262	1:	Y11970	:	Y02934	1:	Y03782		Y03794	LELVLLKA		w97840 c
Ck: 1679 len: 207 ! Breast cancer associated antigen precursor <(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q x(158)(A)X(S)(P)XX(I)(T)(H)XX(P)XXX(P)X(33) MSPLLRRLLLAALLQLAPAQAPVSQPDAPGHQRKVVSWIDVYTRATCQPREVVVPLTVELMGTVAN MSPLLRRLLLAALLQLAPAQAPVSQPDAPGHQRKVVSWIDVYTRATCQPREVVVPLTVELMGTVAN	**(*) 1 (*) 7 (*)	ck: 5919 len: 243 Wheat Type III glutathione transferase subu <(x)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C)(P,Y,S)(P,Y,Z)(P,Y,Z)(P,X,Z)(Z)(Z)(Z)(Z)(Z)(Z)(Z)(Z)(Z)(Z)(Z)(Z)(Z	{1,200 80}(V) OFPES	ck: 4119 len: 102 ! Human 5' EST secreted protein SEQ ID No: 59){1,200}(L,I,V,M,A,P)X(P,T)(P)x(P)(E)xx(V)(S)(K)xx(M) PKPPKPVSKMRMATPLLMQALPM	ck. 3125 len: 24 ! Human CLIP immunomodulatory peptide. Use of	MASMNSKOPHRAMHFILFERHA SABARARARA 1: GGGGPGGGLLGGSAHPHPHMHSLJGHLSHPAAAAANNEPSGLEHPGLVAAAAHHGAA AAAAAAAGOVAAASAAAAVVGAAGLASICDSDTDPRELEAFAERFKQRRIKLGVTQADVGSALANLKIPGVGSLSQSTICRFESLTL AAAAAAAAGOVAAASAAAAVVGAAGLASICDSDTDPRELEAFAERFKQRRIKLGVTQADVGSALANLKIPGVGSLSQSTICRFESLTL	<pre><(x)(1,200)(L,1,V,M,A,P)X(P,T,S)(L,1,V,M,A,P)XX(L,1,V,M,A,F,Y,W)(Q) <(x)(1,200)(L,Xx(L)(S)(H)XX(A)XXX(M)X(211) x(196)(M)X(S)(L)XX(L)(S)(H)XX(A)XXX(D)XX(D)XX(L)XX(D)XX(D)XX(D)XX(D)XX(</pre>	ck: 3324 len: 423 ! Brn-3a polypeptide. New polypeptide compris	<pre><(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(Q x(9)(V)x(P)(L)xx(L)(S)(R)xx(V)xxx(A)x[88] x(9)(X)x(P)(L)xx(L)(S)(R)xx(V)xxx(A)x[88] xx(A)x(B) xx(A)x(B)x(A,B) xx(A)x(B)x(B)x(B)x(B)x(B)x(B)x(B)x(B)x(B)x(B</pre>	ck: 1817 len: 113 ! Human 5' EST secreted protein SEQ ID No: 57	<pre><(X){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,E,I,W)(Y) x(3){A)x(S)(P)xx(L)(S)(R)xx(M)xxx(C)x(34) TLVAGSPCSLSRWIMAGFCHGELVQSDMESQEWERGQVVLSHTSLPWCYVSPR</pre>	1 12	<pre></pre> <pre><</pre>	ck: 881 len: 135 ! S. aureus polypeptide. New essential polype	YNKGPQPIQRIRDRV	Ck: 7277 len: 144 ! S. aureus polypeptide: New Gascillar Folif (Ck: 7277 len: 144 ! S. aureus polypeptide: New Gascillar Folif (Ck: 7277 len: 144 S. aureus polypeptide: New Gascillar Folif	(SETASSH	1 (x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W)(C,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y	ck: 7697 len: 207 Winged bean chymotrypsin inhibitor WCI-3. A
در مر		4	н		щ			р .		1 DLRQSLARALKVN GALRFAGNLLIHR		RALKVNARPLTVAE NLLIHRAARRRVLI	ц		GWRLPEYTLSQEGG ISPENHISLTNVVG	4		Ľ	.	AAPSTTSALTPGPAA
	Y07241	Y07240	 H	Y07239	∺	Y04743	. 1	Y 04 9 0 1		l: arpltvaev aarrkvllm	Y04960	MRHTFRRH		Y04959	PAHKREYT' HSLGCTWH		W93953 (H	707007 c	aaaadaaassvakgga
<pre>x(X)(1,200)(b,1,7,7,7,7,7,7,7,7,7,7,7,7,7,7,7,7,7,7,</pre>		ESSKEGG	EGYIMGFISKERERAILSTKPPG	ck: 2144 len: 141 ! Fragment of human praceurer state Fragment of human pr	(S)(R)xx(V)xxx(A)x(80) ASRYLVVPRATLATWADGYERRPANAP	ck: 5429 len: 110 ! Mycobacterium species protein sequence c	ISKAT	1,200 L.T.Y.M.A.P)X(P,T,S)(L,T,V,M,A,P)XX(L,T,V,M,A,P)XX(L,T,V,M,A,P)XX(L,T,V,M,A,P)XX(A)XXX(S)X(S)	ck. 6286 len: 72 Mycobacterium species protein sequence 3	L (192) (P) X (T) (Y) XX (N) (S) (H) XX (A) XXX (L) X (72) (P) X (T) (Y) X (M) (S) (H) XX (A) XXX (L) X (72) (P) X (T) X	ck: 8320 len: 280 ! Mycobacterium Species Figure A.F.Y.V.	IGIVVEHVVTASMALLOPRRVIADEGLLGALKERG IRERLTAVAIVAHKPHVDS	<pre>(X)[1,200](L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,I,V,M,A,F,F,I,T,M,A,F,F,I,T,M,A,F,I,X,I,Y,X,I,X,I</pre>	ck: 8454 len: 273 ! Mycobacterium species protein sequence 4	TICRLESEMETOKGASKKOAKRNAAEKFLAKESN SLRNSPGEKINLLKRSLLSIPNTDYIQLLSEIAKEQGFNITYLDIDELSANGQYQCLAELSTS	<pre><(x){1,200}(L,I,V,M,A,P)X(P,T,S)(L,I,V,M,A,P)XX(L,I,V,M,A,F,Y,W,X,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y,Y</pre>	ck: 7759 len: 352 ! Human regulatory molecule HRM-9 protein.	X(4/)(V)X(3)(r)XX(2)(V)XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX	I'A'W'	

Databases

searched:

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TILIAN ON THE PROPERTY OF THE 
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Eppstein-Barr virus; EBV; EBV &
Epstein-Barr virus
Epstein-Barr virus
                                                                                                                                                                                                                 17-MAY-1989.
10-NOY-1988; 310591.
11-NOY-1988; IE-003041.
(GGEN-) G Gene Galway Ltd.
Gannon BFX, Walls DFG;
WPI; 89-146591/20.
N-PSDB; N91751.
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Total length: 23,686,106
:al sequences: 188,963
CPU time: 03:36.13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            _SEQUENCE 1.0
P91884 standard; peptide; 48
P91884;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New immunoglobulin E competitor
- binding exclusively with low affinity receptors (FcER2)
Disclosure; Table 4; 33pp; English.
Amino acid residues (see corresp. DNA N90237) representing
mouse gamma-2b chain. Used to extend p90006 (as Region_2)
create an IgE competitor binding to type 2 low affinity
Fc receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          _SEQUENCE 1.0 P90009 standard; protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (RECO) Research Corp Ltd.
Gould HJ, Helm BA, Marsh PJHB;
WPI: 89-178361/24.
N-PSDB; N90237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mouse gamma-2b chain 
Immunoglobulin E competitor;
                                                                                                                            Identifying antigens from a pathogenic organism by generating a gene bank using nucleic acid fragments and screening with infected serum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bennich enumeration; reduced
                        Claim 8(3); 21pp; English.
It is coded for by a DNA sequence isolated by using sera
infected with EBV to screen colonies of micro-organisms t
fragments of EBV DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                51
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              receptors.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length: 102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PVLDSDGSYF IYSKLDIKTS KWEKTDSFSC NVRHEGLKNY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PQVYILPPPA EQLSRKDVSL TCLVVGFNPG DISVEWTSNG HTEENYKDTA
48 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                antigen;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   vaccine;
                                                                                                                                                                                        expressed
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                                                      a from patients
transformed with
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DT AC

_SEQUENCE 1.0
P94157 standard; protein;
P94157;
P94157;
12-JUN-1990 (first entry)

246

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P91884

Length: 48 February 11,

2000 15:48

Type: P

Check:

718

RGWFCPSLCP

SEEPGTSGTP

EPLGPASRRP

PGLRSPLSPV KPKECLRG

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!!AA_SEQUENCE 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P94157
                                                                                                                                                                                   P90460
                                                                                                                                                                                                                     DNA encoding snake venom fibrolase
- for use as thrombolytic agent.
Disclosure; fig 8; 41pp; English.
Alpha-factor profibrolase fusion construct (see N90243) of
plasmid pxS308. Used as a thrombolytic agent to treat
vertebrates, esp. humans. Produced by recombinant techniques.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Constructing new polyester biopolymers - using genes encoding beta-ketothiolase(s), acetoacetyl-CoA reductase(s) and polyhydroxybutyrate synthetase(s).

Disclosure; p; English.

The enzyme is used to study the polyhydroxybutyrate (PHB) biosynthetic pathway. It catalyses the reduction of acetoacetyl CoA to form D(-)-beta-hydroxybutyryl-COA, the substrate for PHB synthetase, and can be used to control biopolymer syntheses and produce new biopolymers. The sequence was deduced from the DNA sequence.

See also p94154-p94156.
                                                                                                                                                                                                                                                                                                                                    12-JUL-1998: 311924.
16-DEC-1988: 311924.
18-DEC-1987: US-134981.
(CHIR) Chiron Corp.
Valenzuela P. Brake A, Randolph
WPI: 89-200623/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-JUN-1988; U02227
29-JUN-1987; US-067695.
29-JUN-1987; US-067695.
(MASI) Massachusetts Institute
Peoples OP, Sinskey AJ;
PEI, 89-039555/05.
N-PSDB; N91209.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Acetyl-CoA reductase.
Polyester biopolymers; acetyl-CoA reductase; polyhydroxybutyrate;
Alcaligenes eutrophus H16.
W08900202-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Alpha-factor profibrolase from Southern Copperhead venom Southern copperhead fibrolase; snake venom fibrolase; alpha-factor profibrolase fusion construct; Crotalidus; thrombolytic agents; plasmid pKS308.

Agkistrodon contortrix contortrix
                                                        151
                                                                                       101
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                                                                                                                                                                                                               vertebrates, esp.
Sequence 288 AA;
                                                                                                                                                                                                                                                                                                                          N-PSDB; N90243
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1-NOV-1989 (first entry)
Alpha-factor profibrolase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          P90460 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                51
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                                                                                                                                                                                 Length: 288
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MTQRIAYVTG GMGGIGTAIC QRLAKDGFRV VAGCGPNSPR REKWLEQQKA
VMAAMLSDQP
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                             GMCQLKHSTG
                                                                                         DHRMNTKYNG
                                                                                                                       AVLPESNSTN NGLLFINTTI
                                                                                                                                                    MRFPSIFTAV LFAASSALAA
                                                          ITVISVSHDT
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                             VIQDHSAINL LVALTMAHEL GHNLGMNHDG
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                                                          LASFGNWRET
SKLFSDCSKK
                                                                                       DSDKIRQWVH
                                                                                                                                                                               February 11,
DYQTFLTVNN
                                                                                                                       ASIAAKEEGV SLDKRQQRFP QRYVQLVIVA
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                                                                                       QIVNTINEIY
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                                                          DLLRRQRHDN
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                                                                                                                                                    AQIPAEAVIG YLDLEGDFDV
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                               NQCHCGANSC
                                                          GDTVGLAYVG
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Inducting immune response with complex of epitope contg molecule and rotavirus VP6 inner capsid protein as carrier, attached and rotavirus VP6 inner capsid protein as carrier, attached without chemical coupling agent

SC claim 9; Page 22; 30pp; English.

This peptide is found to bind to rotavirus protein VP6. It is considered from rotavirus VP3 protein. Ability to bind is due to considered from rotavirus VP3 protein. Ability to bind is due to considered from rotavirus VP3 protein. Ability to bind is due to considered from rotavirus VP3 protein and an Arg in the 3-D protein. The spatial arrangement of a Cys and an Arg in the 3-D protein. The spatial arrangement of a compsn for inducing immunological coupled to an epitope-bearing molecule via a binding sequence. Coupled to an epitope-bearing molecule via a binding sequence. See also P81997 and P81999-P82000).

See also P81997 and P81999-P82000).

Gee also P81997 and P81999-P82000) and avoids converse effects or antigenicity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P81999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Binding peptide 2 capable of b rotavirus strain S-All; VP6 in gastroenteritis; ss.
P70477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE 1.0
                                                                                                                                                                            LJ-LEC-1986; U02756.

14-JAN-1986; US-818740.
(UYNC-) UNIV OF WAROLINA.
(WERT/) WERTZ G W.
(WERT/) WERTZ G W.
WPI; 87-206300/29.
N-PSDB; N70785.
Vaccines fo-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYSA-) Univ Saskatchewan.
Sabara M I, Frenchick P J, Mullin-Ready K F;
WPI; 88-055927/10.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE 1.0
                                                                                             fragment encoded by a DNA sequence of fragment encoded by a DNA sequence of syncytial virus proteins.

Syncytial virus proteins.

Disclosure; Chart 14; 57pp; English.

A novel plasmid which comprises a DNA novel plasmid which comprises a DNA protein, and the protein itself, are
                                                                                                                                                                                                                                                                                    yuman respiratory syncytial virus (HRSV).
408704185-A.
16-JUL-1987.
13-DEC-1986; U02756.
                                                                                                                                                                                                                                                                                                                                                                                       5-APR-1991 (firs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length: 25
                                                                                                                                                                                                                                                                                                                                                                          protein.
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Length: 194
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                                                                                                                                                                                                                                                                                                                                                                                                                                         standard;
                                               can
                                          plasmid which comprises a DNA sequence encoding this and the protein itself, are claimed, for use as HRSV. The vaccine can be administered to pregnant women or child bearing age to stimulate maternal antibodies. can also be vaccinated at 2-3 months of age.
                                  n also be
                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
numan respiratory
                                                                                                                                                             human respiratory virus - comprising proteins or 
pded by a DNA sequence coding for human respiratory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          February 11, 2000 15:48
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                                                                                                                                                                                                                                                                                                                                                                                                                                           protein; 194
   February
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       11,
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                                                                                                                                                                                                                                                                                                                                                                                               syncytial virus
       2000 15:48
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     Type: P Check: 9828
                                                                                                                                                                                                                                                                                                                                                                                                     (HRSV)
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g peptide;
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MSRRNPCKFE
DKSIDTLSEI
KLLTELNSDD

IRGHCLNGKR SGAAELDRTE

CHFSHNYFEW EYALGVVGVL NSPKIRVYNT

PPHALLVRQN

ESYIGSINNI

TKQSACVAMS

VISYIESNRK NNKOTIHLLK

R24296 Length: 384

February

11,

2000

15:48

Type:

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| IAA_SEQUENCE 1.0 | ID p70785 | standa AC p70785; | DT 05-APR-1991 | DE 22K protein. | KW Vaccine; ss. | OS Human respitator | PN 16-JUL-1987; | PF 23-DEC-1986; | PR (UTMC-) UTMC-) UTMC-) | OTM | PT Vaccines | PR | OTM | PT Vaccines | PT | PSDB; | P7041 | PT Vaccines | PT | PSDB; | P7041 | PT Vaccines | PT | PSDB; | PTOTE | PT | PSDB; | PTOTE | PSDB; | PTOTE | PSDB; | PTOTE | PT
      !!AA_SEQUENCE 1.0
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polypeptides involved in expression of glycopeptide antibiotic presistance - useful in diagnosing presence of Gram-positive resistance - useful in diagnosing presence of Gram-positive resistance - useful in diagnosing presence of Gram-positive resistance strains e.g. Enterococcus faecium and E gallinarum from the plasmid prise from the ded., 319:157-161). The vans protein is part of a two component (regulatory system for controlling the amount of expression of the regulatory system for proteins VanH, VanA and VanX. The other component of the regulatory system is the VanR transcription regulator. VanS protein can modify phosphorylation of VanR in transcription. See also Q251/8-Q25183.
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(UYNC-) UNIV OF N CAROLINA.
(WERT/) WERTZ G W.
(WERT/) WERTZ G W.
WERT/) WERTZ G W.
WPI, 87-206300/29.
p-PSDB; P70477.
Vaccines for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vaccines for human respiratory virus - comprising proteins or fragment encoded by a DNA sequence coding for human respiratory syncytial virus proteins.

pisclosure; Chart 14; 57pp; English, pisclosure; Chart 14; 57pp; English, are claimed, for use as HRSV protein, and the protein itself, are claimed, for use as HRSV rotein, and the protein tiself, are claimed, pregnant women or vaccines. The vaccine can be administered to pregnant women or child bearing age to stimulate maternal antibodies. Infants can also be vaccinated at 2-3 months of age. Sequence 194 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-OCT-1990; FR-013579.
(INSP) INST PASTEUR.
AITHUM M, COUTVAlta
WDT.
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20-NOV-1992 (first entry)
Regulatory protein Vans involved in glycopeptide resistance.
Regulatory protein vancomycin; telooplanin; resistant;
Clycopeptide antiblotic; vancomycin; tensposon;
clycopeptide antiblotic; peptidoglycan precursor; transposon;
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equence encoding human respiratory syncytial virus (HRSV) A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 inverted repeats.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ur M, Courvalin P, Dutka-malen S, Molinas C; 92-183677/22.
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AEQRKNDVVM

YLAHDIKTPL TSIIGYLSLL DEAPDMPVDQ

LCRVMLSKFA LEKREQDAKL KAKYVHITLD

101

SILENKYDLN
KYFDEINTGI

DVLIQNEDKQ

SIRNNIDIFI IELSAEMDVM

EQKLNTLKRT

HLDAMKLYQY

YVAIVISILI

51

LVIKLKNKKN DYSKLERKLY MYIVAIVVVA IVFVLYIRSM IRGKLGDWIL

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ID AAC DT DE KW
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A_SEQUENCE 1.0

R24190 standard; Protein; 186

R24190;
24-NOV-1992 (first entry)
Bovine RSV strain A 51908 M2 p:
Bovine respiratory syncytial v:
M2 gene; BRSV.
Bovine respiratory syncytial v:
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N-PSDB; 01895.

N-PSDB; 01895.

N-PSDB; 1aevis bone morphogenetic protein and DNA encoding it used in therapy of fracture or osteoporosis

used in therapy of fracture or osteoporosis

claim 2; Fig 4; 28pp; English.

A Xenopus laevis unfertilised egg cDNA library in lambda gtl0 was screened with a PstI-HindIII fragment of X.laevis Xarl4 chromosomal screened with a PstI-HindIII fragment of X.laevis Xarl4 chromosomal DNA. Three clones were isolated, including clone Xbr22 which was found to encode a protein having homology with X.laevis BMP-2A. They were subcloned in pUC19 and used to transform competent E.coli HB101 cells. Transformant E.coli HB101/pxbr22 coding for the BMP-2A was sequenced and the amino acid sequence of BMP-2A deduced from it.

See also Q10890-4 and Q10896-7.

commence 348 AA:
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Xenopus Bone Morphogenetic Factor BMP-2A.
BMF; osteoporosis; fracture; cartilage.
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WPI; 91-075112/11.
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06-SEP-1989; JP-229250.
20-JUL-1990; JP-190774.
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 virus
                           protein.
virus; vaccine; diagnosis;
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strain A 51908
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                              antibodies;
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Rhinovirus. EP-261403-A.

30-MAR-1988. 20-AUG-1987; 112104 P81093; 21-MAR-1991 (first entry) Sequence of rhinovirus HRV2 viral protein Passive immunity; diagnosis; therapy; ss.

P81093 standard;

protein; 323

51

GVTDYIHMLG IIIRNSSDPQ

EAFGNGFVDS TILATLTLIG

VKEHIHAINP CSGSPWRFLK

VGNISKKIIK WMLRIISAMV EKFCKWTQLN YIHKE P81092

Length:

95

February 11,

2000 15:48

Type: P

Check: 3060

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14-MAY-1992.
04-NOV-1991;
05-NOV-1990;
                  DNA molecules corresponding to all or part of the RNA of rhinovirus strain HRV89 (Fig 4, N81390) is claimed, esp. the portion encoding the viral proteins VPI-VP4, P2A-P2C, P3A-P3C. Also claimed are the polypeptides encoded by any of these DNA molecules. The polypeptides are used for stimulating a protective immune response and for blocking cellular receptors. Ab are useful for assay and for purifice. of the corresp. antigen, and can also be used for the therapeutic and diagnostic applications.
                                                                                                                                                                                                                                                                                                                                                                                                               EP-261403-A.
30-MAR-1988.
20-AUG-1987; 112104.
17-JAN-1987; DE-701301.
(BOEH) Boehringer Insommergruber W, Neubauer C, Ruchler E, Frasel L, Zorn M;
WPI; 88-085735/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bovine respiratory syncytial virus genes - used in the prodn. of agents for use in detection and as vaccines for BRSV infection. Claim 32; Page 57; 74pp; English.

This is the sequence of bovine respiratory syncytial virus (BRSV) strain AS1908 M2 protein. It can be used in the detection of BRSV antibodies and in vaccines to prevent infection. It can also be used for the production of BRSV protein antibodies.

See also R24184-R24191 and R25310.
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Sequence of rhinovirus HRV2 viral
Passive immunity; diagnosis; there
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WPI; 92-183675/22.
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                                                                                                                                                                                                                                                                                           Example; Fig 20; 66pp; German.
                                                                                                                                                                                                                                                                                                                                                     New DNA corresponding to viral RNA of rhino-virus HRV89 - useful prodn. of polypeptide(s) for stimulating immune system against HF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P81092 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Rhinovirus
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US-608937
       8
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TNYSTAKAGL

HGFTMALAGE VATKGVTVNT VSPGYIATDM VKAIRODVLD

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!!AA_SEQUENCE 1.0
ID R10974 standard; Protein; 246
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Duechler M, Skern T, Sommergru Kuchler E, Frasei L, Zorn M; WPI; 88-065735/13.
                                                                                    R10974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example; Fig 20; 66pp; German.

DNA molecules corresponding to all or part of the RNA of rhinovirus strain HRV9 (Fig 4, N81390) is claimed, esp. the portion encoding the viral proteins Vpl-Vp4, P2A-P2C, P3A-P3C. Also claimed are the polypeptides encoded by any of these DNA molecules. The polypeptides are used for stimulating a protective immune response and for allocking cellular receptors. Ab are useful for assay and for purificen of the corresp. antigen, and can also be used for the therapeutic and diagnostic applications.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-JAN-1987; DE-701301
(BOEH) Boehringer Inge
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                                                                                                                                                       bacteria or plants, acetyl CoA reductase is an enzyme pisclosure; fig 3; 64pp; English.

This Alcaligenes eutrophus acetyl CoA reductase is an enzyme which is essential to the biosynthesis of polyhydroxyalkanoate which is essential to the biosynthesia of plasmid clone, (PHA). The gene encoding this is contained in plasmid clone, (PHA), downstream from the thiolase gene. The use of recombinant methods for producing such enzymes, required for polyester biopolymer synthesis, allows for the control and modification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                      17 APR-1991 (first entry)
Acetoacetyl CoA reductase enzyme.
Polyester biopolymers; polyhydroxybutyrate; polyhydroxy
beta-ketothiolase; acetoacetyl CoA reductase.
                                                                                                                                                                                                                                                                    Construction and modification of polyester bio:polymers .introduction of poly-hydroxy-butyrate and -alkanoate genes
                                                                                                                                                                                                                                                                                                                      (MASI ) MASSACHUSETTS I
Peoples OP, Sinskey AJ;
WPI; 91-051341/07
                                                                                                              synthesis process.
See also Q10499-500 and Q10502-03.
Sequence 246 AA;
                                                                                                                                                                                                                                                                                                                                                             0-JUL-1990;
101
                                                                                                                                                                                                                                                                                                                                                                                                                      lcaligenes eutrophus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 KLNLYEKQVE SLRAVDMKTQ EKIKMEIDTL
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                                                                                    Length:
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TRADWDAVID TNLTSLFNVT KQVIDGMADR
                                                        MTQRIAYVTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            corresponding to viral RNA of rhino-virus HRV89
of polypeptide(s) for stimulating immune system a
                           LGYDFIASEG
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                               NVADWDSTKT
                                                            GMGGIGTAIC
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                               AFDKVKSEVG
                                                            QRLAKDGFRV
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                                  EVDVLINNAG
      GWGRIVNISS
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                                     ITRDVVFRKM
                                                                   REKWLEQQKP
                                                                                             Check: 3836
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s into
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| ! AA_SEQUENCE 1.0
| ID R12468 stand
| AC R12468;
| DT 29-AUG-1910
| DE Luffa cylind
| OS Luffa cylind
| OS Luffa cylind
| PN J0310938-A
| PD 09-MAY-1991
| PF 22-SEP-1989
| PA (20MO) SUMO
| DR WEI; 91-81-
| PT New protein
| PT Inhibiting |
| PS Claim 1; Fil
| PS Claim 1; Fil
| PS Claim 1; Fil
| CC It has prote
| CC see also J0
| SQ Sequence
                 I AA_SEQUENCE
ID R13252 st
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SUMO ) SUMITOMO CHEM IND WPI; 91-181458/25.
                                                                                                                                                                                               WO9110685-A.
25-JUL-1991.
17-JAN-1991;
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               The CCP-1 coding sequence was isolated from the amino acid sequence of CCP-1 was predicted from the CDA sequence of the Sequence was solated from the cytotoxic T-cell Title CRP-1 coding sequence was isolated from the cytotoxic T-cell Pile MTL 2.8.2 generated from CDA sequence. The structure of the protein was used to design peptides which competitively inhibit the protease. See also 012863-6 and RI3254-RI3262.
                                                                                                                                                                                                                                                                                                            11-OCT-1991 (first entry)
Murine Cytotoxic Cell Protease-1.
mouse; CCP-1 inhibitor; cytotoxic T-lymphocytes;
                                                                                                                                                                                                                                                  protein
                                                                                                                                                                                                                                                                          peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      51
                                                                                                                                                                                                                                                                                                Mus musculus
                                                                                                                                                                19-JAN-1990; US-467880.
(SERA-) SERAGEN INC.
Bleackley RC, Lobe CG, Paetkau VH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEP-1989;
                                                                                                                                                    leackley RC, Lobe
oi; 91-237989/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length: 248 February 11,
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                                                                                                                                                                                                                                                                                                                                                              standard;
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                                                                                                                                                                                                       U00340
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                                                                                                                                                                                                                                                                                                                                                                                                                           SRFKYIEGQI
                                                                                                                                                                                                                                        /label= signal_peptide
21. .247
/label= CCP-1
                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                             Protein; 247
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                                                                                                                                                                       James
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Type: P
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    Check: 528
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R13252

Length:

247

Febřuary 11,

2000 15:48

SRPYMALLSI

KDQQPEAICG

MKILLLLTL SLASRTKAGE IIGGHEVKPH

APIGKYSNTL

LLKLKSKAKR TRAVRPLNLP

RRNVNVKPGD

GFLIREDFVL

TAAHCEGSII

NVTLGAHNIK EQEKTQQVIP

MVKSIPHPDY VCYVAGWGRM

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III.
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                                                                             R11247 standard: Protein; 14
R14247;
03-JAN-1992 (first entry)
HCV-T (6110-6557) encoded ep
Epitope; PCR; diagnosis.
W05114779-A.
W0511479-A.
W0511479-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 _SEQUENCE 1
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06-MAR-1985; US-709183.
07-MAR-1985; US-709183.
(WHITT-) WHITEHEAD INST BIOM.
Alper S1, Kopito RR;
WPI: 86-25231/38.
WP-PSDB; N66551.
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Disclosure; Fig. 9; 49pp; English.
Band 3 is the major glycoprotein of mammalian erythrocytes and functions as an anion antiport in that it mediates the 1-for-1 exchange of chloride (Cl.) and bicarbonate (HCO3-) across the erythrocyte plasma membrane. The anion transport protein may be used in the detection of genetic defects resulting in abnormal transport of ions across cell membranes, e.g. as in cystic fibrosis. N60550 and p60644.
ITR.) MITSUI TOATSU CHEM INC.
kada T, Enomoto N, Date T, Nakao T;
T; 91-310579/42.
PSDB; Q14076.
W nucleotide sequences encoding HCV epitope(s) - f
hepatitis C virus infection via polymerase chain
sclosure; Fig 1(A)-(B); 79pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P60645;
10-MAR-1993
01-JAN-1980
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01-JAN-1980 (first entry)
Mouse kidney cell Band 3-like anion transport protein.
Band 3; anion transport protein; glycoprotein; cystic disease diagnosis; erythrocyte; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mus musculus.
WO8605203-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                VDYSIEDTYT QKLSVPSGFS VTAPDKRGWV INPLGEKTPF PVWMMVASLL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; Protein; 149
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                                                                                                                                                                                                                                                                                                                                                                                                       epitope
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12-DEC-1991.
31-MAY 1991; G00869.
01-UUN-1990; GB-012196.
(ISIS:) ISIS INNOVATION LT
Wakefield AE, Hopkin JM, M
WPI; 92-007487/01.

LID.

ER;

01-APR-1992 (first entry)
NADH dehydrogenase subunit 2
Pneumonia; assay; AIDS; immun
Pneumocystis carinii.
W09119005-A.
12-DEC-1991.

immunosuppressed N

N-PSDB; 020065.

New DNA sequences which act as oligo:nucleotide primers - assaying DNA sample from respiratory secretion of a patien infected with p carinii
Claim 3; Fig 3; 42pp; English.
The amino acid sequence is that of P. carinii NADH dehydro abonit 2 which was translated from DNA from plasmid pAZI: Sequence 482 AA;

carinii NADH dehydrogenase NA from plasmid pAZ112. See

patient for ID R21409 stand

standard; Protein;

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IIAA_SEQUENCE 1.0

ID R20004 standar

AC R20004

DE VP6 inner caps

KW Rotavirus; vac

KW Rotavirus; vac

KW ROTAVIRUS; vac

FN USSO71651-A.

PD 10 DEC-191.

PF 05-MAR-1990; (
PR 02-SEP-1986; [
PR 02-SEP-1987; [
PR 03-SEP-1987; [
PR 03-SEP-1987; [
PR 05-MAR-1990; 
                                                                                                                                                             AC R20004; standard; peptide; 25 AA.

AC R20004;

DT 25-MAR-1992 (first entry)

DE VP6 inner capsid epitope, 84 TS, Peptide B.

KW kotavirus; vaccine; gastroenteritis; virus protein 6; hapten;

PN 05-MAR-1990; 489790.

PN 05-MAR-1990; 489790.

PR 03-SEP-1987; US-092120.

PR 03-SEP-1987; US-092120.

PR 03-SEP-1987; US-092120.

PR 03-SEP-1987; US-092120.

PR 03-SEP-1980; US-092120.

PR 03-MAR-1990; US-082120.

PR 03-SEP-1980; US-092120.

PR 03-SEP-1980; US-092120.
                                                                                                                R20004 Length:
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1 CNIAPASIVS RNIVYTRAQP NQDIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 51
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                                                                                               15:48
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R21409

Length: 482

MLLSSIISQL

IAIALSSSHW NLVLLSRISI ISLIYSIILT

February 11, 2000 15:48

Type:

ש

Check: 617

YNYYYVEIIG

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IIAA_SEQUENCE 1.0
IID R22695; standard; Protein; 286 Aa R26695; 1992 (first entry)
DI 10-NOV-1992 (first entry)
DE blaC31 mutation (5).
Beta-lactanase; BapMI excision 1
KW antibiotic; transformants.
OS Escherichia coll.
DCGation/Qualifi.
FH Key
FT misc_difference 194. .198
FT w09207090-A.
                                                                                                                                                                                                                                                                                                                                                                                             !!AA_SEQUENCE 1.0
                                                                                                                                                                                                                                                                                                                                                               R23004 Length: 297 February 11,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-MAR-1992.
24-JUL-1990; 194069.
24-JUL-1990; JP-194069.
(KIBU-) KIBÜN KK.
(KIBU-) KIBÜN KK.
(KITA) KITASATO RES INST.
WPI; 92-137918/17.
                                                                                                                                                                                                                                                                                                                                                                                      New virulence plasmid pKDSC50 - provides DNA fragment useful as probe for detecting Salmonella Disclosure; Fig 2; 9pp; Japanese.

The 32 kD protein may be transcribed from the first reading frame of the "mba" fragment of virulent plasmid pKDSC50 from nucleotide 1159-2052. Three other proteins may also be transcribed from the Salmonella "mba" fragment, which is capable of causing mouse bacteremia. See also R23005-7.

Sequence 297 AA;
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Protein transcribed from the mba sequence of plasmid Virulent; Salmonella Choleraesuis; mouse bacteremia; Salmonella choleraesuis.
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                                                                                                                                                                                                                                                                                                                                    MDFLINKKLK IFITLMETGS FSIATSVLYI TRTPLSRVIS DLERELKQRL
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                                                                                                                                                                                                                                                   FHRESLYCRT SVEGGVMLFI PKKFFLCGKP
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                                                                          ise; BspMI excision linker; human
transformants.
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                              Location/Qualifiers
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                                                                                         hormone;
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II AA_SEQUENCE 1.0
IID R25601
AC R25601
AC R25601
DE NANH 1993
DE NANH 1913
DE NANH 1913
DE NANH 1913
DE NANH 1913
PN J04169193-A.
PP 19-0CT 1990;
PR (YAMA) YAMAN
DR WPI 92-25339
PR NT1GENIC POLITIFIC POLIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Provided to determine effect of modified DNA prilibraries - used to determine effect of modification on interaction of encoded product, e.g. enzyme with target printeraction of encoded product, e.g. enzyme with that the product which is come of a group of inkers designated black].

CC within the scope of the invention, this sequence was used to illustrate a novel method of modifying DNA. DNA libraries certain the scope of the invention product and be used to determine the susceptibility of a particular crelationship between structure and function. Modified beta-lactamse can be used to determine the susceptibility of a particular cantibiotic to hydrolysis le. neutralization by mutant enzymes. Since call possible combinations of mutations were represented within the scope cof the invention, a meaningful sample could be achieved. The method of contain a mutation linker, to determine the number of different colonies present in the library. This can then be used to calculate the present that the most and the least common codon combinations are present to detect and anlyse structure/function relationships.
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21-OCT-1991, U5-602158.
22-OCT-1990, U5-602158.
(GETH) GENEWTECH INC.
BOTSTEAD D, PALZKILL T,
WPI, 92-167168/20.
Antigenic polypeptide of non-A, non-B hepatitis virus - for diagnosing non-A hepatitis virus infections, also used as vaccine claim 7; Page 2; 10pp; Japanese.

The sequence given is an antigenic polypeptide of non-A, non-B hepatitis (NANBH) virus. This polypeptide is useful for the detection of NANBH virus antibodies. This antigen can detect the virus antibodies in both the acute phase and chronic phase. The detection of NANBH virus genome using this antigen is very sensitive. Polyclonal and monoclonal antibodies for the detection of the virus antigen and the medical treatment of NANBH, can be prepared using the antigenic peptide. This peptide can be used in the production of a vaccine. Sequence 85 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NANBH virus antigen.

Antibody; acute phase; chronic phase;
Non-A, non-B hepatitis virus.

J04169193-A.

17-JUN-1992.
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                                                                                                                                                                                                                                                                                                                                                                                                                      (YAMA ) YAMANOUCHI PHARM CO LTD. WPI; 92-253396/31.
N-PSDB; Q26616.
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09-OCT-1990; JP-272980
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R25601

Length:

85

February 11,

2000 15:48

Type:

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Check: 5700

KKVTFDRLQV LDDHYRDVLK EMKAKASTVK AKLLSVEEAC KLTPPHSAKS

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!!AA_SEQUENCE 1.0
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26-JUN-1992.
11-NOV-1990; J04417.
11-NOV-1990; JF-304417.
(TOKU) TOKUYAMA SODA KK.
WPI; 92-263663/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis C virus antigen expressed as recombinant in E.coli-
useful for diagnosis of hepatitis C virus infection
Claim 1; Page 5; 65pp; Japanese.
The sequences given in R25854-74 are hepatitis C virus protiens.
The genes encoding these proteins can each be used to prepare
recombinant vectors by ligating the gene of interest in to a vector
to be expressed in E. coli. These polypeptides are useful as
diagnostic reagents for type C hepatitis and they may be produced
efficiently by recombinant methods.

Sequence 239 AA;
         Hepatitis C virus antigen expressed as recombinant in E.coliuseful for diagnosis of hepatitis C virus infection Disclosure: Fig 17: 66pp. Japanese.

The sequences given in R25876-95 are encoded by the claimed hepatitis C virus genes of the invention which have been inserted into an E. colivector. These polypeptides are useful as diagnostic reagents for type C hepatitis and they may be produced efficiently by recombinant DNA techniques.
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HCV polypeptide 16.
Recombinant vector; E. coli; diagnostic; reagent; type C hepatitis.
Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   201
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Sequence
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Hepatitis C
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J04179482-A.
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WPI; 92-263663/32.
                                                                                                                                                   N-PSDB; Q27018.
                                                                                                                                                                                                                                                                                                                                                                                                                 R25891 standard; Protein; 253
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                                                                                                                                                                                                                                                                                                                                  vector; E. coli; diagnostic; reagent;
virus.
                                                                                                                                                                                                                                                            /*tag= a
/note= "Sequence R25869"
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                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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R25891

Length: 253

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MITPSLHACM QVVCCSMSYT WTGALITPCA AEESKLPINP LSNSLLRHHS

MITSUBISHI KASEI

February 11, 2000 15:48 Type: P Check: 4644

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ID R25303 standard; Protein; 194
!!AA_SEQUENCE 1.0
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14-JAN-1986; US-818740.
13-JUL-1988; US-218737.
(UYNC-) UNIV NORTH CAROLINA.
COLLINS D., Westz GW;
WPI; 92-340247/41.
         11-JUN-1992; 109812.
11-JUN-1991; JP-139268.
11-JUN-1991; JP-179794.
07-OCT-1991; JP-277008.
16-DEC-1991; JP-33329.
20-APR-1992; JP-099957.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 19; 21pp; English.

The sequences of mRNA encoding HRSV structural proteins are given in Q29627-26. The proteins are F, G, 22K, 9.5K and major capsid protein N. The sequences and encoded proteins are useful for preparing vaccines against HRSV. The vaccines can be used to confer immunity against respiratory tract infections on human subjects. Sequence 194 AA;
                                                                                                                                                                                                       26-APR-1993 (first entry)
HCV NS4-NS5 peptide NZ9-1, NZ9-2, NZ9-3.
Clone; polypeptide; NS4-NS5; Hepatitis C; Virus; HCV;
transcriptase; cDNA; primer; allele; core; region; ups
hydrophilic; turn structure; alpha helix; beta sheet;
determinant; antiserum.
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16-DEC-1992.
                                                                                                                                      misc_difference
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Vaccines for human respiratory virus - include structural genes coding for native structural viral proteins and immunogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HRSV 22K protein.
Vaccine; human respiratory syncytial virus; HRSV; major capsid protein; N.
Human respiratory syncytial virus strain A2.
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                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                determinant; anti
Hepatitis C virus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length: 194 February 11, 2000 15:48 Type: P Check: 9668
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pp 16-DEC-1992; 109812.

pp 11-JUN-1992; 109812.

pr 11-JUN-1991; JP-139268.

pr 11-JUN-1991; JP-139268.

pr 20-APR-1992; JP-28708.

pr 20-APR-1992; JP-33329.

pr 20-APR-1992; JP-039357.

pr (MITU) MITSUBISH KASEI CORP.

pr Hayashi N, Honda Y, Murakami T, Seki M, Takahashi K;

pr Teranishi Y;

pr Wari 92-41721/51.

DR N-psDB; 032510.

RN-psDB; 032510.

CR Hayashi N, Honda Y, Murakami T, Seki M, Takahashi K;

pr diagnosing and vaccinating against hepatitis C virus infections

pr diagnosing and vaccinating against hepatitis C virus infections

pr diagnosing and vaccinating against hepatitis C virus infections

pr diagnosing and vaccinating against hepatitis C virus infections

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pr diagnosing and vaccinating against hepatitis C virus infections

pr diagnosing and vaccinating and are highly reactive with the core region of the Hov gene of the thought to acc as antigen determinants and are highly reactive with accinations and are highly reactive with the core also 032435.
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DR NPSDB; 032505. virus gene and its encoded protein - used for PT New hepatitis C virus gene and its encoded protein - used for PT New hepatitis C virus infections protein of the protein vaccinating against hepatitis C virus infections protein superiors given in R29811-906 and R29533 are encoded by various C clones of the NS4-NS5 regions of the Hepatitis C Virus (HCV) gene of C the invention. These NS4-NS5 RNA sequencess were isolated from the C the invention. These NS4-NS5 RNA sequences were isolated RNA serum of a patient suffering from hepatitis C (HC). The isolated RNA cerum of a patient suffering from hepatitis C (HC). The isolated RNA cerum of a patient suffering from hepatitis C (HC). The sequence cofficient of the sequences were converted into CDNA using transcriptase in the presence then amplified using primer pairs. The CDNA sequences isolated C represent different alleles of the same region of the HCV gene.

C contain a highly higher pairs. The converse thought to sequence analysis shows that these clones represent the core region which is not an alpha helix or a beta sheet. These polypeptides are thought to act as antigen determinants and are highly reactive with C antisecrum raised against HCV-associated antigens. See also Q32436.
!!AA_SEQUENCE 1.0
ID R29879 stand
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Teranishi Y
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26-APR-1993 (first entry)
HCV NS4-NS5 peptide 2918.
Clone; polypeptide; NS4-NS5; Hepatitis C; Virus; HCV; serum; HC;
transcriptase; cDNA; primer; allele; core; region; upstream;
hydrophilic; turn structure; alpha helix; beta sheet; antigen;
determinant; antiserum.
Hepatitis C virus
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EP-518313
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E DAGE

26-APR-1993 (first entry) HCV NS4-NS5 peptide N29-2. _SEQUENCE 1. R29896 star

standard; Protein; 171

101

DLGVRVCEKM ALYDVVSTLP QAVMGSSYGF QYSPGQRVEF LVNAWKSKKS

51

KDLLEDTETP

IDTTIMAKNE VFCVQPEKGG RKPARLIVFP

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ID R29895 stan
                                                               Well 92-417213/51.

DR N-PSDB; 033526.

Virus gene and its encoded protein - used for property of the partitis C virus infections of the partitis C virus (HCV) gene of the sequences given in R29871-906 and R29533 are encoded by various of the invention. These NS4-NS5 regions of the Hepartitis C (HC) which is colated from the clones of the NS4-NS5 regions of the Hepartitis C (HC). The isolated RNA of the invention of a patient sequences were isolated from the crown of a patient sequences given in 032565-77. The sequences were converted into cDNA using transcriptase in the presence of one of the primer sequences given in 032565-77. The sequences were contain amplified using primer pairs. The CDNA sequences isolated trapscriptase in different alleles of the same region of the HCV gene.

Contain a highly hydrophilic region which can adopt a furn structure contains a highly hydrophilic region which can adopt a furn structure with thought to act as antigen determinants and are highly reactive with antigen determinants and are highly reactive with can attagence 171 AA;
                                    R29895
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11-JUN-1991; 109812.
11-JUN-1991; JP-139268.
12-JUL-1991; JP-172794.
07-0CT-1991; JP-287008.
16-DEC-1991; JP-33239.
20-APR-1992; JP-33239.
(MITU) MITSUBISHI KASEI CORP.
(MITU) MITSUBISHI KASEI CORP.
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26-ApR-1993 (first entry)
4CV NS4-NS5 peptide N29-1.
Clone; polypeptide; NS4-NS5; Hepatitis C; Virus; HCV; serum; HC;
Clone; polypeptide; NS4-NS5; Hepatitis C; Virus; HCV; serum; HC;
transcriptase; cDNA; primer; allele; core; region; upstream;
hydrophilic; turn structure; alpha helix; beta sheet; antigen;
hydrophilic; turn structure; alpha helix; beta sheet; antigen;
determinant; antiserum.
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                                      Length: 171
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YRDVLKEMKA KASTVKAKLL
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                                      February 11,
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    SVEEACKLTP
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                                        2000 15:48 Type: P
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    PHSARSKFGY
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The Counce; polypeptide N29-3;
We Clone; polypeptide; NS4-NS5; Hepatitis C; Virus; HCV; serum; HC;
We With transcriptase; cDNA; primer; allele; core; region; upstream;
We with the content of the core; antigen; turn structure; alpha helix; beta sheet; antigen; with the core; antigen; antige
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PA (MITU) MITSUBISHI KASEI CORP.
PI Hayashi N, Honda Y, Murakami T, Seki M, Takahashi K;
PI Teranishi Y;
PI New hepatitis C virus gene and its encoded protein - used for diagnosing and vaccinating against hepatitis C virus infections protein respectively.
PI New hepatitis C virus gene and R29533 are encoded by various convention. These NA4-NS5 regions of the Hepatitis C virus (HCV) gene of the invention. These NA4-NS5 RNA sequences were isolated from the sequences were converted into cDNA using transcriptase in the presence of one of the primer sequences given in 032565-77. The isolated RNA company of the amplified using primer pairs. The cDNA sequences were then amplified using primer pairs. The cDNA sequences region contain an highly hydrophilic region which can adopt a "turn structure" which is not an alpha helix or a beta sheet. These polypeptides are thought to act as antigen determinants and are highly reactive with cantiserum ratsed against HCV-associated antigens. See also Q32436.
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determinant; antiserum.
Hepatitis C virus
EP-518313-A.
16-DEC-1992; 109812.
11-JUN-1992; 109812.
11-JUN-1991; JP-139268.
12-JUL-1991; JP-172794.
07-OCT-1991; JP-277008.
16-DEC-1991; JP-332329.
20-ABR-1992; JP-099957.
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R29897 standa
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transcriptase; cDNA; primer;
hydrophilic; turn structure;
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III.AA. SEQUENCE 1.0
IID R29910 standard; F
AC R29910;
DT 05-MAY-1993 (firs
DE Prod. of the luffi
NKW Loofah; amplificat

101

VYNITLLLSS ASGASRYTLM TLSNYDGKAI TVAVDVTNVY IMGYLVNSTS MKRFTVLILA IFVAASTVEA DVRFSLSGSS STSYSKFIGD LRKALPSNGT

ASQYVFKGST IVTLPYSGNY

EKLQTAAGKI

YDSTAAAAF LVIIQTTAEA SRFKYIEGQI IERISKNQVP

PVVITDDKGQ

201 151

VVTKNIQLLL NYKQNVAAFD

EDVSAKE AQTINGTEKT

R29910; 05-MAY-1993 (first entry) Prod. of the luffin-g gene. Loofah; amplification; missile treatment;

Protein;

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IIAA_SEQUENCE 1.0

ID R29909 standard; Protein; 277 AF
AC R29909;
DT R29909 of the luffin-f gene.

NEW Loofah; amplification; missile to Loufah cylindrica.

PN J04320687-A.

PN J04320687-A.

PN J04320687-B.

PN (NISB) JAPAN TOBACCO INC.

DR WPI; 92-426681/52.

DR N-PSDB; Q32687-B.

PT Viral diseases in agriculture province of the sequence is that of the province of the sequence is the province of the sequence is the province of the province of the sequence of the seq
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                                                                      PD 11-NOV-1992.

PD 18-APR-1991; 086857.

PR 18-APR-1991; DP-086857.

PR (NISB) JAPAN TOBACCO INC.

PR (NISB) JAPAN TOBACCO INC.

PR NP1; 92-426681/52.

DR NP1; 92-426681/52.

DR N-PSDB; Q32887.

PT New gene coding luffin - for missile treatment and prevention of private diseases in agriculture

PT New gene coding luffin - for missile treatment and prevention of private diseases in agriculture

PT New gene coding luffin - for missile treatment and prevention of the luffin fene of Luffa cDNA coding a propertion of the produce was obtd. by PCR of luffa cDNA coding a PCR primer designed based on the luffin protein sequence coding a PCR primer designed based on the luffin gene can be used in coding a PCR primer designed based on the luffin gene can be used in the luffin may be produced stably, by transforming a microbe such as E. coli coding a propertion of the luffin gene.

CC Sec also R23910.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The sequences given in R29871-906 and R29533 are encoded by various clones of the NS4-NS5 regions of the Hepatitis C Virus (HCV) gene of the invention. These NS4-NS5 RNA sequencess were isolated from the serum of a patient suffering from hepatitis C (HC). The isolated RNA sequences were converted into cDNA using transcriptase in the presence of one of the primer sequences given in Q32565-77. The sequences were then amplified using primer pairs. The CDNA sequences were represent different alleles of the same region of the HCV gene. Sequence analysis shows that these clones represent the core region contain a highly hydrophilic region which can adopt a "turn structure" which is not an alpha helix or a beta sheet. These polypeptides are thought to act as antisen determinants and are highly reactive with antiserum raised against HCV-associated antisens. See also Q32436.
     R29909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       R29897 Length: 167 February 11, 2000 15:48 Type: P Check: 2886
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05-MAX-1993 (first entry)
Drodd. of the luffin-f gene.
Loofah; amplification; missile treatment; virus; agriculture.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             101
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Length: 277 February 11, 2000 15:48 Type: P
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!!AA_SEQUENCE 1.0
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See also R22903.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-APR-1991; 086857.
18-APR-1991; JP-086857.
(NISB ) JAPAN TOBACCO INC.
WPI; 92-426681/52.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Luffa cylindrica
J04320687-A.
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                                             Nucleotide sequences hybridising to regions of chicken nov gene useful as probes for detecting complementary sequences to evaluate development and/or differentiation of tumours claim 21; Page 39; 67pp; French.

The chicken nov gene is stimulated in avian nephroblastoma induced pavian myeloblastoma virus but not in normal adult kidney. A laysbe conducted control of the chicken embryonic fibroblasts screened with a tumour-derived from chicken embryonic fibroblasts screened with a tumour-derived from chicken embryonic fibroblasts screened with a tumour-derived probe. Fragment XXII (036044) is part of the 3rd and 4th exons of the nov gene; nucleotide sequences which hybridise to Fragment XXII the nov gene; nucleotide sequences which hybridise to Fragment XXII (181608) or an amino acid sequence 60% homologous to it.
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R31608;
24-MAY-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                              stringent hybridisation;
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                                                                                                                                                                                                                                                                                             25-JUN-1992; F00589.
25-JUN-1991; FR-007807.
(CNRS ) CENT NAT RECH SCI.
Martinerie C, Perbal B;
WPI; 93-036377/04.
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Length:
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    205
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  February 11,
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      2000 15:48 Type: P
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          Check: 7012
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VCDEPKDQTV

TCLDGAVGCM LGPASRVSRV

FLXVRVVILS

SPDCPFPRRV QGGSPNCADR

KLPGKCCEEW

PDTEWSACSK

DVRVPQCLTS

ASPTPLFPSS PLCSMDVRLP

SPAKDGAPCI FGGTVYRSGE

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II AA SEQUENCE

II R30618 st
AC R30618 r
AC R30619 p
AC CLUCK-) L
AC R0F2 r
AC CLUCK-) L
AC R0F2 r
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                 IL R32192 standard; Protein; 246
pr Transgenic plants producing poly:hydroxy-alkanoate polymer(s) -
pr obtd. by transformation with DNA encoding 3-ketothiolase,
pr obtd. by transformation with DNA encoding 3-ketothiolase
pr obtd. by transformation with DNA encoding a peptide which exhibits 3-ketothilase activity,
competity DNA encoding a peptide which exhibits 3-ketothilase activity,
competity DNA encoding a peptide which exhibits 3-ketothilase activity,
competity DNA encoding a peptide which exhibits 3-ketothilase activity,
competity DNA encoding a peptide which exhibits 3-ketothilase activity,
competity DNA encoding a peptide which exhibits 3-ketothilase activity,
competitive encoding enc
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19-MAY-1993
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(UYYMA-) UNIV MADISON JAMES.
(UNMS ) UNIV MICHIGAN STATE.
Dennis DE, Politier Y, Some
WHI: 93-058785/07.
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tho JM, Choi DY, Kim CH, Kim ST, Lee YB, Lim KJ, Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAY-1993 (first entry)
30-MAY-1993 (first entry)
Sequence encoded by the acetoacetyl-CoA reductase (phbb) gene
polyhydroxybutyrate (pHB) operon
Operon; polyhydroxyalkanoate; acetoacetyl-CoA reductase.
Alcaligenes eutrophus.
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KR-009510.
KR-013601.
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R32192

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PD 13 MAY-1993; U09487.

PD 13 MAY-1993; U09487.

PR 04-NOV-1991; US-787567.

PR 04-NOV-1992; US-901707.

PR 19-JUN-1992; US-901707.

PR 19-JU
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                08 JUL-1993.
30-DEC-1992; U1327.
31-DEC-1991; US-816283.
(ARCH-) ARCH DEV CORP.
Bell GI, Scino S, Yamac
WPI: 93-22727/28.
WP-PSDB; Q45653.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        an antibody Iray...
248 AA;
                                                                                                                                                                                                                                                                                                                                                                                R39259 standard;
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Plant type I RIP Luffin.
Type I ribosome-inactivating protein; ricin; gelonin; momordin;
immunoconjugate; autoimmune disease; cell killing; toxin; loofah.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE 1.0
                                                                                                                                                                                                                                  18-NOV-1993 (first entry)
18-NOV-1993 (first entry)
Human somatostatin receptor-1.
Human; somatostatin; receptor; SSTR-1; SSTR-2; SSTR-3; tumour;
pancreas; 1slet; promoter; transformation; host cell.
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                                                                                                                                                                                                      Homo sapiens.
WO9313130-A.
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  Somatostatin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   7294 standard; protein; 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length: 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length: 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                               LAQTNNGTFK TPVVITDDKG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MTQRIAYVTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IVTLPYSGNY EKLQTAAGKI REKIPLGFPA LDSALTTIFH YDSTAAAAAF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DVRFSLSGSS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            fragment.
  receptors useful for
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SRFKYIEGQI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STSYSKFIGD LRKALPSNGT VYNLTILLSS ASGASRYTLM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                February 11, 2000 15:48
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                                                                                                                                                                                                                                                                                                                                                                                Protein;
                                                                               Yamada
                                                                                                                                                                                                                                                                                                                                                                                   391
                                                                                                                                                                                                                                                                                                                                                                                                                                                            QRVEITNVTS
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  diagnosis of tumours -
                                                                                                                                                                                                                                                                                                                                                                                                                                                            KVVTKNIQLL LNYKQNVA
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!!AA_SEQUENCE 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R39260;
18-NOV-1993 (first entry)
Murine somatostatin receptor-1.
Mouse; somatostatin; receptor; SSTR-1; SSTR-2; SSTR-3;
pancreas; islet; promoter; transformation; host cell.
Mus.musculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 3: Page 63-64; 94pp; English.

The sequences given in R39259, R39251 and R39253 represent the human somatostatin receptors (SSTR)-1, SSTR-2 and SSTR-3. The DNA encoding these proteins was isolated from total human pancreatic islet DNA. These DNA sequences may be placed under the control of a suitable promoter and used to transform a host cell. The DNA sequences and these proteins may be used in screening assays for testing candidates including agonists and antagonists of SSTR polypeptides. The assays may be used to discriminate candidate substances with desirable properties specific to SSTR polypeptides. The isolated substances may be used in a wide range of aplications eg. diagnosis of various human tumours. Fragments of these DNA sequences may be used as probes in the isolation of other SSTR-encoding clones.
                    Claim 3; Page 65-66; 94pp; English.

The sequences given in R39260, R39262 and R39264 represent the murine somatostatin receptors (SSTR)-1, SSTR-2 and SSTR-3. The DNA encoding these proteins was isolated from total murine pancreatic islet DNA. These DNA sequences may be placed under the control of a suitable promoter and used to transform a host cell. The DNA sequences and these proteins may be used in screening assays for testing candidates including agonists and antagonists of SSTR polypeptides. The assays may be used to discriminate candidate substances with desirable properties specific to SSTR polypeptides. The isolated substances properties specific to SSTR polypeptides. The isolated substances properties specific to SSTR polypeptides. The isolated substances human tumours. Fragments of these DNA sequences may be used as 201 % and 201 % a
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Bell GI, Seino S, Ya
WPI; 93-227272/28.
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30-DEC-1992; US-816283.
31-DEC-1991; US-816283.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Somatostatin receptors useful for diagnosis of tumours - also useful for screening candidate somatostatin receptor agonists
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Yamada Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  391
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R39260

Length: 391

February 11,

2000 15:48 Type: P

Check: 8110

MFPNGTASSP SSSPSPSPGS CGEGACSRGP GSGAADGMEE PGRNASQNGT

ISFIYSVVCL

VGLCGNSMVI YVILRYAKMK TATNIYILNL

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!!AA_SEQUENCE 1
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                                                                                                                                                                                                                                                                                                                                                     peptide(5) and nucleic acid sequences related to EBV - for detecting EBV in samples Claim 3; Page 40; 62pp; English. Claim 3; Page 40; 62pp; English. The sequence is that of Epstein Barr Virus viral capsid antigen VCA-p40 encoded by the EBV open reading frame BdRFI. This antigen is immunochemically reactive with antibodies to EBV and can be used for the diagnosis of EBV in a sample. See also R42316-35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Epstein-barr virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Viral capsid antigen;
                                                                                                                                                                                                                                             151
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                                                                                                                                                                                                     251
                                                17-FEB-1994.
06-AUG-1993;
07-AUG-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2-MAR-1993;
3-MAR-1992;
                 Cochran
WPI; 94
                                                                             Equine herpesvirus (Strain: Dutta;
WO9403628-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 iddeldorp JM;
PI; 93-345368/44.
                                                                                                            Equine herpesvirus;
                                                                                                             -AUG-1994 (first entry)
line herpesvirus US2 gene.
line herpesvirus; US2; vac
                                                                                                                                                                                                                                                                                                                          Length: 345 February 11, 2000 15:48
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recombinant equine
                                                                                                                                                                                  RGKKLVQASA
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        94-065715/08.
DB; Q56616.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; Protein;
                                                                                                                                                   standard; Protein; 303
                                       1993; U07424.
1992; US-926784.
SYNTRO CORP.
                                                                                                                                                                                                                                                                                                                                                  345 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry) from ORF BdRF1.
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EP-200721.
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   herpes
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                                                                                                             vaccine; antigen; protection;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Barr Virus;
                                                                                                                                                      A
                                                                                           Individual
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                                                                                                                                                                                                         RNSGPREDTN
                                                                                                                                                                                                                            QSYEVPRYVP
                                                                                                                                                                                                                                                AGLSKSVNEL
                                                                                                                                                                                                                                                                    PAPPAACSHT
     used
                                                                                                                                                                                                                                                                                                                               Type: P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             viral genome; antibodies;
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                                                                                              isolate: S-1EHV-000)
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     prepare
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                                                                                                                                                                                                          PQQPTTEGHE
                                                                                                                                                                                                                              HPPPPPTSHC
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                                                                                                                  prophylaxis;
        vaccines
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DE TO A

R49297; 16-SEP-1994 Invarient chain

(first entry) nain (li) posit

position

97-120

standard;

Protein; 24

R49296

Length: 25

February 11,

2000

15:48

Type: P

Check: 4901

LPKPPKPVSK MRMATPLLMQ

ALPMG

Sequence

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ILAA_SEQUENCE 1.0
ID R49296 standard; Protein;
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11-AUG-1993; US-925460.
11-AUG-1993; US-925460.
15-JUN-1993; US-925460.
15-JUN-1993; US-925460.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to protect horses from infectious equine herpes virus Example 1; Page 83-84; 153pp; English.

Recombinant equine herpesvirus' (EHV) have a foreign DNA sequence inserted into their genomes. The inserted sequence is a piece of foreign, double stranded DNA which encodes an RNA molecule which foreign double occur in the animal into which the EHV is does not naturally occur in the animal into which the EHV is introduced. The foreign DNA sequence is introduced into the EHV virus a homology vector. The recombinant EHV's are used to prepare a homology vector horses from infectious EHV. This is the US2 gene product from a recombinant EHV. Recombinant attenuated EHV's sequence 303 AA;
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                                              Disclosure: Page 36; 139pp; English. R46981-7038 represent peptide The sequences given in R49291-505 and R46981-7038 represent peptide fragments of naturally-occuring immunomodulatory proteins. These fragments are between 10-30 residues in length and bind to a human fragments are between 10-30 residues in length and bind to a human major histocompatibility complex (MHC) class II allotype. These peptides may be used for therapy of autoinmune diseases, such as peptides may be used for therapy of autoinmune diseases, and to type I diabetes, rheumatoid arthritis and mutiple sclerosis, and to type I diabetes, rheumatoid arthritis and mutiple sclerosis, and to reduce transplant rejection. They may also be used for vaccination providing an exclusively T-cell-mediated response, which can be class I or class II based, or both, depending on the length and length and length and length and length and length are length and length are length and leng
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16-SEP-1994 (first entry)
Invarient chain (11) position 97-121.
Naturally-occuring; immunomodulatory protein; human; therapmaturally-occompatibility complex; class II; ellotype; type major histocompatibility complex; class II; cell-mediated autoimmune disease; rheumatoid arthritis; Tecl-ore; MHC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          mutiple sclerosis; transplant rejection;
                                                                                                                                                                                                                                                                                                                                              Novel immunomodulatory peptide(s) treatment of auto:immune diseases,
                                                                                                                                                                                                                                                                                                                                                                                                                                Vignali
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chicz RM,
                            character of the immunogenic peptides.
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Wovel immunomodulatory peptide(s) and nucleic acids - useful for treatment of auto:immune diseases, transplant rejection and for vaccination

Disclosure; Page 36; 139pp; English.

Disclosure; Page 36; 139pp; English.

The sequences given in R49291-505 and R46981-7038 represent peptide

fragments of naturally-occuring immunomodulatory proteins. These

fragments are between 10-30 residues in length and bind to a human

major histocompatibility complex (MHC) class II allotype. These

peptides may be used for therapy of autoimmune diseases, such as

type I diabetes, rheumatoid arthritis and mutiple sclerosis, and to

reduce transplant rejection. They may also be used for vaccination

providing an exclusively T-cell-mediated response, which can be

class I or class-II based, or both, depending on the length and

commance 34 Ma.
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Homo sapiens.

W09404171-A.
03-MAR-1994.
11-AUG-1993; U07545.
11-AUG-1992; US-925460.
15-JUN-1993; US-925460.
(HARD ) HARVARD COLLEGE.
(HARD ) Hedley ML, Stern LJ, Strominger
                                                                                                                                                                                                                                                                                                                               16-SEP-1994 (first entry)
Invarient chain (11) position 97-119.
Invarient chain (11) position 97-119.
Naturally-occuring; immunomodulatory protein; human; therapy; class I; major histocompatibility complex; class II; allotype; type I diabetes; autoimmune disease; rheumatoid arthritis; T-cell-mediated response; mutiple sclerosis; transplant rejection; vaccine; MHC.
Homo sapiens.
M99404171-A.
103-MAR-1994.
11-AUG-1993; U07-92460.
11-AUG-1993; U07-92460.
15-JUN-1993; US-922460.
15-JUN-1993; US-922460.
15-JUN-1993; US-922460.
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Chicz RM, Hedley
Vignali DA;
WPI; 94-082825/10.
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R49299 standa
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IIAA_SEQUENCE 1.0

IID R49438 standard; PAC R49438;

DE R49438;

DE Minigene HLA-DRalp EMINIGENE HLA-DRALP MATURALLY-occuring KW Maturally-occuring KW major histocompati KW autoimmune disease KW mutiple sclerosis; PN W09404171-A.

PD 11-AUG-1993; U07-94

PF 11-AUG-1993; U07-94

PF 11-AUG-1992; U57-92

PR 11-AUG-1992; U57-92

PR 11-AUG-1993; U57-92

PR 11-AUG-1992; R49438 AA_SEQUENCE 1.0
R49589 standard; Protein; 49 AA.
R49589; R49589;
15-SEP-1994 (first entry)
Sequence of HLA-DR alpha chain leader per amino terminus of a blocking peptide fragional in it.
Minigene; HLA-DR alpha leader peptide; ci Homo sapiens.

W09404171-A.

03-MAR-1994.
11-AUG-1993; U07-925460.
11-AUG-1992; US-925460.
15-JUN-1993; US-925460.
(HARD) HARVARD COLLEGE.
Chicz RM, Hedley ML, St
Vignali DA,
WPI; 94-083825/10. WO9404557-A. 03-MAR-1994. 11-AUG-1992; 11-AUG-1992; Disclosure: Page 93; 139pp; English.

The sequences given in R49291-505 and R46981-7038 represent peptide fragments of naturally-occuring immunomodulatory proteins. These fragments are between 10-30 residues in length and bind to a human major histocompatibility complex (MHC) class II allotype. These peptides may be used for therapy of autotimmune diseases, such as type I diabetes, rheumatoid arthritis and mutiple sclerosis, and to reduce transplant rejection. They may also be used for vaccination providing an exclusively T-cell-mediated response, which can be class I or class-II based, or both, depending on the length and character of the immunogenic peptides. 16-SBP-1994 (first entry)
Minigene HLA-DRalpha chain/Ii 24 residue peptide fusion.
Minigene HLA-DRalpha chain/Ii 24 residue peptide fusion.
Naturally-occuring; immunomodulatory protein; human; therapy; class I;
Naturally-occuring; immunomodulatory protein; allotype; type I diabetes;
major histocompatibility complex; class I; allotype; type I diabetes;
autoimmune disease; rheumatoid arthritiss I; -cell-mediated response;
muntiple sclerosis; transplant rejection; vaccine; MHC. Example; Figure 3B; 60pp; English.

The inventors claim a nucleic acid encoding a polypeptide comprising an AA sequence identical to that of a segment of a naturally-occurring human protein which binds to a human major histocompatibility complex (MHC) class II allotype and a second sequence substantially identical to human II in order to construct such minigenes, overlapping synthetic oligos were used to generate the leader peptide/blocking peptide N-PSDB; Q44334. New peptide binding t treating auto-immune (HARD) H Chicz RM, Vignali DAA; WPI; 94-083102/10. peptide Synthetic. Key Novel immunomodulatory peptide(s) and treatment of auto:immune diseases, tra immunisation Length: 49 MAISGVPVLG 1992; U06692. 1992; WO-U06692. HARVARD COLLEGE. M, Hedley ML, Stern 49 AA; February 11, FFIIAVLMSA QESWALPKPP /label= HLA-DR alpha leader peptide 26. .49 /label= chain Ii Location/Qualifiers to MHC class II allotype - useful for e diseases, transplant rejection and for Stern 49 Ę 2000 15:48 Ę Ą Strominger Strominger and nucleic acids - us transplant rejection r peptide linked to the fragment of human invariant KPVSKMRMAT PLLMQALPM chain Type: P Ii. JL, Check: 4300 Urban useful on and : RG for

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The sequences given in R50068 and R50070-82 represent antigens of structural and non-structural regions of non A non B hepatitis virus (NANBHY). The CDNA encoding these sequences wase derived from the plasma of a NANBH patient by recombinant DNA techniques. These fragments are useful for the diagnosis of NANBH patients and the detection of NANBHV carriers.

Sequence 352 AA;
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constructs
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01-JUN-1993; 156087.
10-JUL-1992; JP-207391.
(KOKU-) KOKUSAI SHIYAKU KK.
(SANWA KAGAKU KENKYUSHO CO.
(TOFU ) TONEN CORP.
(TOKR-) ZH TOKYOTO RINSHO IGAKU SOGO KENKYUSHO.
WPI; 94-128677/16.
15-MAR-1994.
15-MAR-1993; 156087.
10-JUL-1993; JF-207391.
10-JUL-1993; JF-207391.
(KOKU-) KOKUSAI SHIYAKU KK.
(SANW) SANWA KAGAKU KENKYUSHO CO.
(TOFU ) TONEN CORP.
(TOKR-) ZH TOKYOTO RINSHO IGAKU SOGO KENKYUSHO.
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NANBH virus antigenic fragment #13.
                                                                                                                                                                                                                             24-NOV-1994 (first entry)
NANBH virus antigenic fragment #14.
Antigen; structural; non-structural; non A
NANBHV; NANBH; patient; plasma; diagnosis;
NOD A NOD B hepatitis virus.
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detection; carrier; ss
                                                                                                                                                                                                                                                                   non B hepa
detection;
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                                                                                                                                                                                                                                                               hepatitis vir
tion; carrier;
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                                                                                                                                                                                                                                                               virus;
ier; ss
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R53646;
18-JAN-1995 (first entry)
c-fos gene product.
c-jun; c-fos; cancer; tumo

vaccine;

; tumour; metastas's; treatment;
lymphocytes; inhibition; antige

antigen;

prophylaxis; n; antitumour

R53646 standard;

Protein;

380

51

GGRVQC MASSMLSSAA R53251

Length:

56

February 11,

2000 15:48

Type: P

VATRINPAQA SMVAPFIGLK SAAFPVSRKQ

NLDITSIASN Check: 2035

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SCCCCCC PPP PR
     !!AA_SEQUENCE 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R50082
protection of glycogen biosynthesis

Disclosure: Page 31, 91pp; English.

The glycogen synthase gene may be incorporated into a plant genome to produce a plant with altered starch synthesising ability. Other genes encoding enzymes involved in the starch or glycogen to biosynthetic pathways may also/additionally be used. The new plants have an improved capacity to produce starch at elevated or lowered temperatures and/or an ability to synthesise starch with an altered fine structure. They can produce higher starch yields at certain temperatures and/or produce starch with an improved quality. The glycogen synthase gene construct used requires the presence of an amyloplast transit peptide to ensure its correct localisation in the amyloplast. This is the consensus sequence of the transit peptide of sequence 56 AA;

Sequence 56 AA;
                                                                                                                                                                                                                                                                         (ZENE ) ZENECA LID
Bird CR, Fentem P
WPI; 94-151328/18.
New plants with all
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 28; Page 32-33; 37pp; Japanese.
The sequences given in R50068 and R50070-82 represent antigens of structural and non-structural regions of non A non B hepatitis virus (NANBHY). The CDNA encoding these sequences wase derived from the plasma of a NANBH patient by recombinant DNA techniques. These fragments are useful for the diagnosis of NANBH patients and the detection of NANBHY carriers.

Sequence 215 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-PSDB; 058827.
Nucleic edid fragment coding non-A non-B hepatitis virus antigen useful in diagnosis of NANB patient and detection of virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        101
                                                                                                                                                                                                                                                                                                                                                                                                                                               Consensus sequence of signal peptide of small subunit of RU Glycogen synthase; starch; gene expression; recombination; transgenic plant; glucose-1-phosphate adenylyl transferase; biosynthesis; metabolism; metabolic pathway.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      201
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length: 215
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                                                                                                                                                                                                                                                             plants with altered starch porating a donor gene spec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard; Protein;
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WO-G01881.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    entry)
                                                                                                                                                                                                                                                                                                           Keeling
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                                                                                                                                                                                                                                                         arch synthesis ability specifying an enzyme
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ion; recombination;
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involved
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pr 28-007-1993; 117519.
pr 28-007-1993; 117519.
pr 28-007-1993; 117519.
pr 29-007-1993; US-968415.
pr 19-007-1993; US-968415.
pr 19-007-1993; US-968415.
pr 19-007-1993; US-968415.
pr 19-007-1993; US-968415.
pr 29-007-1993; US-968415.
pr 
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09-JUN-1994.
17-NOV-1993; U11153.
17-NOV-1992; US-977452.
(ICOS-) ICOS CORP.
GODISKA R, GTAY PW, SC.
WEI; 94-200264/24.
N-PSDB; Q66167.
                                                                                                                                 W09412635-A.
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R53748 standard;
R53748;
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Seven transmembrane receptor (V28).
Primer; seven transmembrane receptor; receptor; amplification;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          351
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EP-599077-A.
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197. 21
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273. .29
/label=
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146. .16
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107. .12
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                                      Schweickart
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                                                                                                                                           Transmembrane
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EP-613946-A.
07-SEP-1994.
11-FEB-1994; 102093.
21-FEB-1994; 102093.
R 21-FEB-1993; JF-024399.
R 28-JUL-1993; JF-186507.
R 28-JUL-1993; JF-186507.
PA (NACA) NAT CANCER CENT.
PA (TAKE) TAKEDA CHEM IND LID.
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activity user its user its user or ulcers of ulcers. Claim 15; page 27; 39pp; English.

This sequence is a heparin-binding secretory transforming factor that sequence is a heparin-binding secretory transforming factor (hst-2) polypeptide. It is preferred that the polypeptides of the invention have regions deleted. The polypeptides have cell growth promoting activity and angiogenesis promoting activity and can be used for the treatment of burns, wounds, postoperative tissues or ulcers. The therapeutic composition can be used as a platelet ulcers. The therapeutic radiotherapy and chemotherapy. (Several contents of the contents of the composition can be used as a platelet composition can be used as a platelet.
                                                                                                                                                                                                                                                                                                                                                                                 misc_difference 2. .30 /note- "may be absent"

/note- "N terminal fragment

EP-613946-A specification"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    R58556 standard; protein; 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-MAR 1995 (first entry)

Reparin-binding secretory transforming factor 2.

heparin-binding growth factor family; fibroblast growth factor;

FGF; hst-2; heparin-binding secretory transforming factor 2;

drugs; reagents; growth promoting activity; increase platelet number;

angiogenesis promoting activity; treatment; burns; wounds; ulcers.

Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 misc_difference
                                                                                                                                                                                           New mutant hst-2 polypeptides - having activity used for treating eg. burns,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 251
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Involving the receptors.

Example 6; Page 65-66; 100pp; English.

Two Synthetic Oligonucleotides (066165, 066166) were annealed and readiolabelled to produce a V28 specific probe. The probe was denatured by boiling for two minutes and then hybridised to the human placenta genomic library. Several strong hybridising signals were identified and plaque purified and the DNA sequence (066167) of a full length genomic clone encoding this seven transmembrane receptor was deduced.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          51
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                                                                                                                                                                                         having cell growth promoting burns, wounds, post:operative
                                                                                                                                                                                                                                                Yoshimura K,
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for conditions
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R58556

Length: 175

February 11,

2000 15:48 Type: P

Check:

IGFHLQVLPD

MLAGEIAGVN WESGYLVGIK RORRLYCNVG

ERGVVSLFGV

RSALFVAMNS KGRLYATPSF

QEECKFRETL

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R63592
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STY2, STY3, STY4 and STY5 cDNAs (given in Q72858-61), encoding peptides having MAP-kinase-phosphatase activity, were derived from human squamous A431 cell polyA+ mRNAs. Residues 177-255 of the encoded STY2-5 peptides were deduced (R63593-96) and demonstrated to show homology with the corresponding region of the previously isolated human CI100 gene product (R63592).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    09-MAY-1995 (first entry)
MAP-kinase-phosphatase CL100 partial sequence.
MAP-kinase-phosphatase; CL100;
MAP-kinase-phosphatase; CL100;
neurological disease; proliferative disorder; inflammation;
Disclosure; Fig. 7d; 53pp; English.
STY8 CDNA (given in Q72864), encoding a peptide having
MAP-Kinase-phosphatase activity, was derived from a human
brain cDNA library. The STY8 protein (R63601) encoded by this
full-length cDNA showed homology to the previously isolated
human CL100 gene peroduct (R63602).
Sequence 367 AA;
                                                                                                                                  13-OCT-1994.
31-MAR 1994; G00694.
31-APR-1993; GB-007250.
10-FEB-1994; GB-007250.
10-FEB-1994; GB-002573.
(CANC-) INST CANCER RES INST.
(CANC-) INST CANCER RES ROYAL CANCER HOSPITAL.
Ashworth A, Hughes DA, Marshall CJ;
WPI; 94-333199/41.
Screening for a substance which is an inhibitor of mammalian kinase - and is used in the mfr. of a medicament for the kinase - and is used in the mfr. of a medicament for the diseasement of e.g. an inflammatory disorder or neurological
                                                                                                                                                                                                                                                                                                                                                                              09-MAY-1995 (first entry)
MAP-Kinase-phosphatase CL100.
MAP-Kinase inhibitor; MAPK; MAP-Kinase-phosphatase; CL100; STY8; neurological disease; proliferative disorder; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Screening for a substance which is an inhibitor of mammalian MAP kinase - and is used in the mfr. of a medicament for the treatment of e.g. an inflammatory disorder or neurological
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(CANC-) INST CANCER RES
Ashworth A, Hughes DA,
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07-APR-1993; GB-007250.
10-FEB-1994; GB-002573.
                                                                                                                                                                                                                                                                                                                                                              cardiovascular disease.
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09-MAY-1995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    standard; Peptide;
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!!AA_SEQUENCE 1.0
ID R58636 standard;
AC R58636;

protein; 124

101

51

SGPWMCYPGY

LDSMYKEHGV QEGQAGTGAF

AFKVPALPGC

RPVLKLQCNG SQVPEAVLRD CCQQLADISE

PSCRREVVKL TAASITAVCK

R58635

Length:

124

February 11,

2000 15:48

Type:

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Claim 1; Page 15; 21pp; English.

CC Inhibitor (AI) family. By size and electrophoretic measurements,

CC inhibitor (AI) family. By size and electrophoretic measurements,

CC this protein has been called 0.26 AI. The protein was isolated from

CC wheat extracts by a novel process comprising (1) extracting a

CC wheat-based material to produce a solution containing the AI; (ii)

CC optionally purifying the solution to remove contaminants using a cation

CC exchanger to absorb the AI, (iii) treating this exchanger with alkali

CC at pH 9-13 to elute the AI, (ii) immediately adjusting the pH of the

CC elute within neutral or acid range and (v) recovering the AI from this

CC solution. The novel AI has a very high inhibitory activity against an

CC amylase especially human pancreatic alpha-amylase. The activity of

CC the AI is 5-30 times higher than other amylases of wheat origin such as

CC 0.53 AI or 0.28 AI. The protein occurs as a homodimer with subunits of

CC molecular masses 12500. The protein can be used in combination with

CC another amylase inhibitor (0.19) (R58656) as an agent for inhibiting

CC suppressing appetite, and as a food additive, the total content of both

CC proteins in the material being at least 20% by weight.
                            TAGGE STANDARD CONTRACTOR OF THE PART OF T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R63602
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05-0CT-1994; 302148.
24-MAR-1994; 3F-091881.
29-MAR-1993; JF-091881.
28-MAX-1993; JF-148423.
(NAGA-) NAGATA SANGYO CO LTD.
(NISS ) NISSHIN FLOUR MILLING CC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              wPI; 94-304407/38.

New amylase inhibitor and process of prepn. from wheat -
useful for inhibiting increase in blood glucose level,
controlling insulin secretion, suppressing appetite and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Amylase inhibitor protein 0.26 AI.

Amylase inhibitor; wheat; cation exchange; inhibitor; amylase;
pancreatic, alpha-amylase; homodiner; blood glucose level;
insulin secretion; appetite; food additive.
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R58636
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CC family. By size and electrophoretic measurements, this protein is called
CO.19 AT. The protein occurs as a homodimer, the subunits having the amino
acid sequence given. This protein can be used in combination with a
convel amylase inhibitor (0.26) (RS8635) as an agent for inhibiting
CI increase in blood glucose level, controlling insulin secretion or
CI suppressing appetite, and as a food additive, the total content of both
CI proteins in the material being at least 20% by weight. The novel protein
CI proteins in the material being at least 20% by weight. The novel protein
CI proteins in the material being at least 20% by weight. The novel protein
CI proteins in the material being at least 20% by meight and containing
CI proteins in the material being at least 20% by meight. The novel protein
CI proteins in the material being at least 20% by meight and containing
CI protein in the axionally purifying the solution to remove contaminants
CI using a cation exchanger to absorb the AI, (ii) treating this exchanger
CI with alkali at pH 9-13 to elute the AI, (ii) immediately adjusting the pH
CI of the elute within neutral or acid arong and (v) recovering the AI from
CI and anylase especially human pancreatic alpha-amylase. The activity of
CI the AI is 5-30 times higher than other amylases of wheat origin such as
CI Sequence 124 as.
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E Amylase inhibitor; protein 0.19 AI.

W Amylase inhibitor; wheat; cation exchange; inhibitor; amylase; hu

W pancreatic; alpha-amylase; homodiner; blood glucose level;

N pancreatic; alpha-amylase; homodiner; homodiner; homodiner; homodiner; homodiner; homodiner; homodiner;

N pancreatic; alpha-amylase; homodiner; 
DNA coding NAD synthetase - used for the prepn. of NAD synthetase Claim 2; Column 17-20; 16pp; Japanese.

The amino acid sequence for the N-terminal region of nicotineamide adenine dinucleotide (NAD) synthetase (NADS). The gene encodes a protein of 245 amino acids. The gene was inserted into E.coli in an expression plasmid. This was used to produce the NADS protein. This method allows the efficient preparation of NADS polypeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      _SEQUENCE 1.0
                                                                                                                                                                                                                                                        01-FEB-1993, 014650.

01-FEB-1993, JP-014650.

(ASAH) ASAHI KASEI KOGYO KK.

WPI, 94-298801/37.

WPSDB: 070537.
                                                                                                                                                                                                                                                                                                                                                                                                          J06225771-A.
16-AUG-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Nicotineamide adenine dinucleotide; Nicotineamide adenine dinucleotide; Baccillus stearothermophilus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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/label=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         synthetase N-terminal f
; NAD; synthetase; NADS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Type: P
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IIAA_SEQUENCE 1.0
ID R63443 standar
AC R63443;
DT O6-JUN-1995
DE Amino terminal
DE fusion enzyme,
Event of the service of the service
NKW enzyme, transg
Synthetic.
PN W09424292-A.
PD 27-OCT-1994;
PR 08-APR-1994;
PR 08-APR-1994;
PR 08-APR-1994;
PR 08-APR-1994;
PR 08-APR-1994;
PR 10-APR-1994;
PR 11-APR-1994;
PR 12-APR-1994;
PR 12-APR-
                   !!AA_SEQUENCE 1.0
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Transgenic starch producing organism - comprises exogenous A glucose pyrophosphorylase enzyme DNA sequence Example; Fig 8; 88pp; English.
Plasmid pPp14 is a pVictorIV SGiN Man deriv. in which a 3.2 EcoRI fragment contg. the construct:
patatin promoter - spinach rubisco activase transit peptide
                                                                                                                                                                                                07-APR-1994; E01082.
08-APR-1993; GB-007408.
(DANI-) DANISCO AS.
Kleczkowski L, Marcussen J,
                                                                                                                                                                                Kleczkowski L,
Villand P;
                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                     Barley; endosperm;
enzyme; transgenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Transgenic starch producing organism - comprises exogenous ADP glucose pyrophosphorylase enzyme DNA sequence Example; Fig 8; 88pp; English.

Plasmid pp54 is a pvictoriv deriv. in which a 3 kb KpnI fragment contg. the construct;

contg. the construct;

patatin promoter - spinach rubisco activase transit peptide - small subunit AGP from barley endosperm - 35S terminator; is inserted in the KpnI site. R63443 gives the N-terminal sequence of the rubisco activase-AGP small subunit fusion enzyme. ppp54 is inserted.
                                                                                                                                                                                                                                                                                                                                                                                                                                             Amino terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                         usion enzyme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R63444 standard; Protein; 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 94-341871/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    fusion enzyme.

Barley; endosperm; ADP glucose pyrophosphorylase; AGP; starch; enzyme; transgenic plant; transit peptide; spinach; pPPS4; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             )8-APR-1993; GB-007408.
(DANI-) DANISCO AS.
Kleczkowski L, Marcussen
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                                                                                                                                                       94-341871/42.
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                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                             ADP glucose pyrophosphorylase; AGP; plant; transit peptide; spinach; pp
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                                                                                                                                                                                                                                                                                                                                                             GP; starch; pPPL4; ss.
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large subunit AGP from barley endosperm - 35s terminator; is inserted in the ECORI site. R63444 gives the N-terminal of the rubisco activase-AGP large subunit fusion enzyme. pF is claimed.

al sequence pppL4

R63444 Length: 120

February 11, 2000 15:48

SSAGASVPTS GFLGSSLKKH TNVRFPSSSR

DSLGIHMQFS

Type: P

Check: 7811

Sequence

120 AA;

101

SVLPLEGKAC VSPVRREGSA

51

TISMIVKAAE NEEKNIDKWA HLAKDFSDDQ LDIRRGKGMV

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MATAVSTYGA ATRAPLNING

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R62507 standa
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15-MAR-1994: AU0126.
16-MAR-1993: AU-007854.
(AUST-) AUGYIN RES INST.
(AUST-) AUGYIN RES INST.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R62507;
26-JUN-1995 (first entry)
26-JUN-1995 (first entry)
7-197tosyl transferase 3' clone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA sequences encoding Gal-alpha (1,3)galactosyl transferase and clones conty. such sequences are used in xenograft therapies Disclosure; Page 31; 50pp; English. The sequence is that of the product of the porcine Gal-alpha (1,3) galactosyl transferase gene which produces a Gal epitope on the surface of porcine cells. This epitope is recognised by antibodies which are responsible for hyperacute rejection of xenotransplanted pig cells, tissues and organs.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Galactosyl transferase 3 (Galactosyl Gal-alpha (1,3) galactosyl
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WO9421799-A.
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WPI; 94-317019/39.
N-PSDB; Q74711.
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                                                                        24-NOV-1994.
18-MAY-1994; U05617.
19-MAY-1993; US-064400.
(ANTI-) ANTIGEN EXPRESS
WPI; 95-006697/01.
Identification of mutant Ii mols
                      Humphreys
WPI; 95-0
                                                                                                                                                                                                                                                                                                                     06355;
08-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                          (66355 standard; peptide;
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                                                                                                                                                                                                                                protein.
protein; immunomodulator;
pro histocompatibility comp
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transferase; xenograft; transplant;
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      having
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      altered
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PROSENTACIO

23-JUN-1995 (first entry)
Human growth differentiation factor-8 protein.
Growth differentiation factor-8; GDF-8; cell p
adipocyte; obesity; transforming growth factor

cell proliferation factor-beta.

Homo sapiens. WO9421681-A. 29-SEP-1994.

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SCCCCPPT
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!!AA_SEQUENCE 1.0
ID R63160 standard; Protein; 375 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cleavage sites - and useful for identifying immunomodulators, esp. peptide(s) which enhance or inhibit antigen presentation bisclosure; Page 38-39; Sopp; English Immunomodulatory peptides of Ii protein (full sequence given Immunomodulatory peptides of Ii protein (full sequence given enhances MHC class II-mediated antigen presentation, while the Ii peptides given in R66353-54 inhibit antigen presentation. Sequence 216 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                23-JUN-1995 (first entry)
Mouse growth differentiation factor-8 protein.
Growth differentiation factor-8; GDP-8; cell proliferation; adipocyte; obesity; transforming growth factor-beta.
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R63159 standa
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                                                                                                                                                                                                                                                                                                  New growth differentiation factor 8 - useful for treatment and diagnosis of cell proliferative disorders esp. of muscle. Claim 3; Page 47; 84pp; English of English be used to maintain cells before transplantation; to improve efficiency of cell fusion and to treat obesity or diseases related to abbormal adipocyte proliferation.
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19-MAR-1993; US-033923.
(UYJO) UNIV JOHNS HOPKINS
Lee S, Mcpherron AC;
WPI; 94-316943/39.
                                          351
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                                            MLYFNGKEQI
                                                                                                                                                                                       DSSDGSLEDD
                                                                                                                                                                                                              YSRIEAIKIQ ILSKLRLETA PNISKDAIRQ LLPRAPPLRE LIDQYDVQRD
                                                                                                                                                                                                                                             MMQKLQMYVY
                                                                       APKRYKANYC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; Protein;
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                                                                                                                                                                                                                                               IYLFMLIAAG PVDLNEGSER EENVEKEGLC
                                                                                                                                                          LRPVKTPTTV
                                            IYGKIPAMVV
                                                                                                     TPKRSRRDFG
                                                                                                                               TVLQNWLKQP
                                                                                                                                                                                                                                                                        February 11, 2000 15:48 Type: P
                                                                          SGECEFVFLQ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               376 AA
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                                                                                                                                                                                        TMPTESDFLM QADGKPKCCF FKFSSKIQYN
                                                                                                                                                             FVQILRLIKP
                                               DRCGCS
                                                                          KYPHTHLVHQ
                                                                                                     LDCDEHSTES
                                                                                                                                    ESNLGIEIKA
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                                                                                                                                    LDENGHDLAV
                                                                                                                                                               MKDGTRYTGI
                                                                            ANPRGSAGPC CTPTKMSPIN
                                                                                                         RCCRYPLTVD FEAFGWDWII
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                                                                                                                                       TFPGPGEDGL
                                                                                                                                                                  RSLKLDMSPG
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IIAA_SEQUENCE 1.0
IID R66665 standar
AC R66655
DT 21-UTL-1995 (
DE Binding peptid
KW Inner capsid |
KW Inner capsid |
KW Dinding peptid
OS Rotavirus
OS OS ROTAVIRUS
OS COUPLED 1987; [
PR 02-SEP-1986; [
PR 03-SEP-1986; [
PR 03-SEP-1987; [
PR 03-SEP-1
                 IIAA_SEQUENCE 1.0
ID R66680 standar
AC R6680;
DT 21-JUL-1995 (
DE Peptide 84 TS-
KW Inner capsid p
KW binding peptid
OS Synthetic.
ON US5374426-A.
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                                                                                                                                                                                                                                                                                    R66665
binding peptide.
synthetic.
US5374426-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-DEC-1994.
03-SEP-1986; 903222.
03-SEP-1986; US-903222.
02-SEP-1987; US-092120.
30-OCT-1989; US-429147.
(UYSA-) UNIV SASKATCHEWAN.
                                                                                                                                                                                                                                                                                                                                           coning molecule.

Claim 11(b): Column 26: 23pp; English.

Claim 11(b): Column 26: 23pp; English.

An immunological carrier complex comprises an epitope-bearing molecule bound to the rotavirus inner capsid protein VP6 (given molecule bound to the rotavirus inner capsid protein VP6 (given molecule bound to the rotavirus inner capsid protein VP6 (given molecule bound to the rotavirus in R66664-8. Reptide B (or 84 TS) (R66665) is derived from rotavirus VP3 (231-254).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-MAR-1994; U03019.
19-MAR-1993; US-033923.
(UYJO ) UNIV JOHNS HOPK.
Lee S, MCPHEITON AC;
WPI; 94-316943/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         351
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                                                                                    Inner capsid prot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunogenic complex contg. rotavirus VP6 protein coupled, esp. by protein-protein interaction, to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Frenchick PJ, Mullin-ready WPI; 95-035623/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  51
                                                                                                                                                                                                                                                                               Length:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LYFNGKEQII
                                                                                                                                                                                                                                       CNIAPASIVS
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capsid pro
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                                                                                                                                                                         standard; peptide;
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                                                                                                                       (first entry)
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                                                                                                                                                                                                                                                                               February 11, 2000
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84 TS)
n; VP6;
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                                                                                    VP6;
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                                                                                                                                                                         25
                                                                                    carrier;
                                                                                                                                                                                                                                       NQDIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sabara
                                                                                                                                                                                                                                                                                  15:48
                                                                                    vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CCRYPLTVDF EAFGWDWIIA
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                                                                                                                                                                                                                                                                                  Type:
                                                                                       rotavirus;
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epitope
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W Synthetic.

Synthetic.

N Synthetic.

N US5374426-A.

PD 20-DEC-1994.

PD 20-DEC-1996; 903222.

PR 03-SEP-1986; 903222.

PR 03-SEP-1989; US-903222.

PR 30-OCT-1989; US-429147.

PA (UYSA-) UNIV SASKATCHEWAN.

PI Frenchick pJ, wallin-ready KF, Sabara MI;

PI Immunogenic complex contg. rotavirus VP6 protein as roupled, esp. by protein-protein interaction, to an optical complex contg.

Provided Protein interaction, to an optical complex contg.

Provided Protein interaction, to an optical complex contg.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PR PR PR PR
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03-SEP-1986; US-903222.
03-SEP-1986; US-903222.
02-SEP-1987; US-092110.
30-OCT-1989; US-092110.
(UYSA-) UNIV SASKATCHEWAN.
Erenchick PJ. Mullin-ready KWPI; 95-035623/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         immunogenic complex contg. rotavirus VP6 protein as carrier coupled, esp. by protein protein interaction, to an epitope contg. molecule.

Example 2; Column 18: 23pp: English.

An immunological carrier complex comprises an epitope-bearing molecule bound to the rotavirus inner capsid protein VP6 (given in R6669) using a binding peptide such as peptide B (84 TS) (R66659) variants of peptide B were also tested for binding to vonly those that generated a spatial arrangement of cys and Argresidues in a suitable 3-dimensional orientation (e.g. R6666-68) were suitable. 84 TS-CYS (R66680) did not bind to VP6.
                                                                       20-JUL-1995 (first entry)
A. aculeatus pectin methylesterase.
Pectin methylesterse; PME; strain CBS 101.43; demet food industry; fruit; vegetable; processing; plant modification; pectin lyase; glucanase; firmness.
Aspergillus aculeatus.
                                                           Aspergillus
WO9425575-A
                                                                                                                                                                                     R67596 standa
                                                                                                                                                                                                                                                                                           An immunological carrier complex comprises an epitope-bearing molecule bound to the rotavirus inner capsid protein VP6 (given in R66669) using a binding peptide such as peptide B (84 TS) (R6665). Variants of peptide B were also tested for binding to VONLY those that generated a spatial arrangement of Cys and Arg residues in a suitable 3-dimensional orientation (e.g. R66666-68) were suitable. MONOSER (R66682) did not bind to VP6. Sequence 25 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE 1.0
                                                                                                                                                     R67596;
20-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         binding
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                                                                                                                                                                                   96 standard;
                                                                                                                                                                                                                                    CNIAPASIVS RNIVYTSAQP NODIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         capsid protein; VP6; carrier; vaccine; rotavirus;
1g peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide;
                                                                                                                                                                                     Protein;
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epitope
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28-APR-1994; DK0173. 30-APR-1993; DK-000487. 28-OCT-1993; DK-001217.

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PI Heldt-Hansen HP, Kauppinen S, Kofod LV;

PI Heldt-Hansen HP, Kauppinen S, Kofod LV;

DR WPI; 95-022253/03.

DR N-PSDB; 079154.

PPI Enzyme with pectin methylesterase activity - useful for price of pectin, e.g. for application in food industry. Price demethylation of pectin, e.g. for application in food industry. Price demethylation of pectin, e.g. for application in food industry.

PRI Enzyme with pectin methylesterase activity - useful for demethylation of general personners an enzyme showing pectin methylesterse (PME) of Strain CBS 101.43. The PME enzyme is useful for demethylation of cotivity. The DNA sequence encoding this protein was isolated from CC strain CBS 101.43. The PME enzyme is useful for demethylation of cotivity. The DNA sequence used in the food industry for fruit and composition. It can therefore be used in the food industry for fruit and vegetable processing. It modifies plant cell walls and can be used to improve the firmness of a pectin-contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. material or to increase viscosity of a pectin-contg. material. CC contg. Material contg. Material contg. Material contg. Material contg. Material contg. Material contg.
                                                       R67596
                                            Transgenic plant material with plastid(s) contg. the enzymes for synthesis of poly:hydroxy:alkanoate(s) - express poly:hydroxy:butyrate and have good growth and seed formation. Claim 2: Page 60-61: 88pp. English.

The acetyl-CoA-reductase gene (phbB) from A. eutrophus is cloned conder the control of an Arabidopsis thaliana seed storage protein conder the control of an Arabidopsis thaliana seed storage protein conderer for plastid tissue-specific gene expression in a conderer for plastid tissue-specifically poly-beta-conderer for plastid tissue-specifically poly-beta-conderer for plastid plastic plant (preferably a Brassica e.g. rape). PHB and related PHAs are conderer for plastid thermoplastics with many useful applications. Sequence 329 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-AUG-1994; U09265;
17-AUG-1993; US-108193.
06-JUN-1994; US-254357.
(UNMS) UNIV MICHIGAN STATE.
Nawrath C. Politier Y, Somer
WPI; 95-098770/13.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (NOVO ) NOVO-NORDISK AS.
Andersen LN, Budolfsen G,
Heldt-Hansen HP, Kauppinen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Acetyl-CoA-reductase; transgenic plant; poly-beta-hydroxybutyrate; biodegradable thermoplastic.
Alcaligenes eutrophus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          301
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-CoA-reductase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; Protein; 329
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February 11, 2000 15:48 Type: P Check: 7564
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R71324

Length: 329

MASMISSSAV TTVSRASRGQ

SAAVAPEGGL KSMTGEPVKK VNTDITSITS

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PAR PRACE
                                                                                                                                         IP R72715 standard;
                                                                                                                                                                                                                                                                                                                                           R70100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          101
                                                                                                                                                                                                                                                                                                                                                                             resistant to virus infection

Claim 1; Fig 14; 92pp; English.

The amino acid sequence encoded by the open reading frame 3 of the real sequence encoded by the open reading frame 3 of the resultant product was amplified by lettuce infectious yellows virus (LIYV) RNA 1. The gene was amplified by PCR using the primers Q83175-6. The resultant product was used to construct expression casesttes expressing the gene as a sense (pEPG170) message. The vectors containing the cassettes can be transformed into marginate the vectors containing the cassettes can be transformed into containing the sense or antisense vectors are resistant to LIYV infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lettuce infectious yellows virus open reading frame 3 protein. Expression vector; expression cassette; Agrobacterium tumefaciens; plant; lettuce infectious yellows virus coat protein; amplify; PCR; primer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     301
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WPI; 95-106554/14.
N-PSDB; 083174.
New DNA encoding proteins of lettuce infectious yellows virus
and related vectors, transformed cells and transgenic plants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  heat shock protein
                                                                                                                                                                                                                                                              101
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                                                                                                                                                                                                                                    151
                                                                                                                                                                                                                                                                                                                                                                   Sequence
                                                                                                                                                                                 251
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                                   WO9510629-A.
20-APR-1995.
                                                                         02-DEC-1995 (first entry)
hisc gene product of Salmonella typhhisc; hisc; Salmonella typhimurium;
identification; mutagen; transition
                                                                                                                                                                                                                                                                                     51
                                                          Salmonella typhimurium.
            14-0CT-1994;
15-0CT-1993;
                                                                                                                                                                                                                                                                                                                                        Length: 274 February 11,
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                                                                                                                                                                                                                                                                                                               MIMMSPLYAL TKQCVIDTAY RLAVPTQHCA
                                                                                                                                                                                    SIVIECGDVI
                                                                                                                                                                                                            LVEKFIFDKL
                                                                                                                                                                                                                                                                                          LCGFKMDTSS
                                                                                                                                                                                                                                                              ELLIGKRLNK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; peptide; 274 AA.
                                                                                                                                                                                                                                                                                                                                                                     274 AA;
 ; U11617.
; US-137627.
V CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Carney KJ,
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                                                                                                                                                                                    GKGINKTKKK
                                                                                                                                                                                                            EFLVYDGEEG
                                                                                                                                                                                                                                     SDKESLNKMN
                                                                                                                                            Protein; 359
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                                                                                                       Salmonella typhimurium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Falk BW,
                                                                                transition; transversion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ISSVNGQKGQ
                                                                                                                                                                                    ICNK
                                                                                                                                                                                                            FKSPQVEYND
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Klaassen V;
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                                                                                               auxotroph;
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PT Salmonella auxotrophic mutants for detecting mutagenic agents - pr comprise point mutation(s) in histidine genes hisG and hisC; for pr identifying specific base changes induced by a mutagen pr identifying specific base changes induced by a mutagen property of the pro
  R72715
US5416202-A.
116-MAY-1995.
09-DEC-1992; 988430.
04-NOV-1991; US-787567.
09-DEC-1992; US-988430.
(XOMA ) XOMA CORP.
Bernhard SL, Better MD, CWPI; 95-193480/25.
                                                                                                                                                                                                                                                                                                                                                                                                                                    misc_difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE 1.0
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                                                                                                                                                                                                                                                                                                                                                                              misc_difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Type I ribosome-inactivating protein luffin. Ribosome inactivating protein; RIP; Type I; cytotoxin;
                                                                                                                                                              misc_difference
                                                                                                                                                                                               misc_difference
                                                                                                                                                                                                                                  misc_difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Plant.
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01-JAN-1996 (first entry)
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: 157. 157
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22
                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                    Carroll SF,
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                    Lane
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R84103

303

February

11,

2000

15:48

Type:

Check: 5723

51

VRSADLYRFS SSLLTLPKAC RPIVRTRGAT AIALDRNGVV YHEDRMGVS

VVDRHKALPN SSIDVDGHLW EFLSRQCFVL ASEPLGIPIV

MGVVLITVVT

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HERE REGERE TO COCCO COC
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CC polypeptides that share the ribosome-inactivating activity of the cc natural protein but differ in AA sequence. Preferred analogues have a Cys available for disulfide bonding located at a posn. It its AA CC sequence from the posn. corresp. to posn. 251 in ricin A-chain RTA CC to the carboxy terminus of the analogue. (R74176 is the sequence of the Type I RIPs gelonin, BRIP, mormordin II, luffin created the Type I RIPs gelonin, BRIP, mormordin II, luffin created in A-chain RTA CC sequence of the Type I RIPs gelonin, BRIP, mormordin II, luffin created in the sequence of the Seq
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16-FEB-1995; U02087.

17-FEB-1994; US-198094.

(SYIR) SYNTRO CORP.

Chiang CH, Cochran MD;
region not essential for replication - used in vaccines to protect horses from infection

Example 11; Pages 86-87; 159pp; English.

T00531 encodes R84103 the equine herpesvirus 1 (EHV) unique short 2
(US2) gene prod. A recombinant attenuated EHV can be produced by deleting the above US2 gene (a region not essential for replication) from the viral DNA. The attenuated virus can be used as a foetal safe vaccine to protect an equine against EHV, or in a test to determine if an equine has been vaccinated against or is infected with EHV.

Sequence 303 AA;
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Key
region
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                                                                                                                                                                                                                                                                                                                                                                                                                                   (SYTR ) SYNTRO COR
Chiang CH, Cochra
WPI; 95-302714/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20-MAR-1996 (first entry)
Equine herpesvirus (EHVI) unique short (US2) gene prod. .
Equine herpesvirus 1; EHVI; unique short; US2; recombinant; vaccine; attenuated virus; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                201
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123. .140
/note= "EHV-1, EHV-4
IBR consereved US2 r.
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265, 8670-74
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I leukocyte adnesion to environment.

T inflammatory disease property in flammatory chain variable in flammatory disease property in flammatory disease antibody 21.6 heavy chain variable region. Ha, directed against leukocyte adhesion molecule via 4. Cloned region, Ha, directed against leukocyte adhesion molecule via constant regions in the construction of a humanized in linked to human constant regions in the construction of a humanized modified using PCR primers (See 099895-98) and then subcloned into modified using PCR primers (See 099895-98) and then subcloned into constant regions. In the humanized light chain, amino acids L45, constant regions. In the humanized light chain, anion acids L45, last and L69 in the humanized light chain, anion acids L45, last anion acid present in the equivalent position of the mouse by the amino acid present in the equivalent position of the mouse cbus the amino acid present in the equivalent position of the mouse by the amino acid present in the equivalent position of the mouse cbus are constant property of the mouse are constant property of the mouse of the mouse constant property of the mous
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Humanized VLA-4 antibody 21.6
Humanized antibody; leukocyte
antibody engineering.
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25-JAN-1995; UO1219.
25-JAN-1994; US-186269.
(APHE-) ATHENA NEUROSCIENCES INC.
Bendig MM, Jones TS, Leger OJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New humanised antibodies
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                                                                                                                                                                                                                                                                           _SEQUENCE 1.0
R81327 standa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            anti-idiotype antibodies.
Sequence 123 AA;
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23-MAR-1996 (first entry)
Mouse VLA-4 antibody 21.6 light heavy variable region.
Humanized antibody; leukocyte adhesion molecule; VLA-4; therapeutic;
                                                                                                                                                                                                                                                                                                                                                                               51
                   region
                                                           region
                                                                                                   peptide
                                                                                                                                           Mus musculus
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    /note=
50. .54
/note=
                                                                                                        Location/Qualifiers
1. .19
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e= "framework region 1"
                                                                                    "signal peptide"
       "complementarity determining region
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The sequence represents the mouse antibody 21.6 heavy chain variable CC region directed against leukocyte adhasion molecule VLA-4. Cloned CC cDNA sequences of mouse 21.6 VH and VL (see 69989) regions are constant regions in the construction of a humanized CC antibody against VLA-4. The 5 and 3 ends of the mouse cDNAs are CC modified using PCR primers (See 099895 98) and then subcloned into CC constant regions. In the humanized heavy chain, amino acids H27. CC constant regions. In the humanized heavy chain, amino acids H27. CC constant regions to the humanized heavy chain, amino acids H27. CC constant regions. The humanized heavy chain, amino acids H27. CC constant regions. The humanized heavy chain and could the mouse could be the amino acid present in the equivalent position of the mouse CC lass. H29, H30, H44 nd H71 in the equivalent position of the mouse CC lass into COS cells. The humanized antibodies can be used to inhibit adhesion of a leukocyte to an endothelial cell and to treat inflammatory diseases such as multiple sclerosis. They can also be used in the treatment of stroke, cerebral traumas, CC can also be used in the treatment of stroke, cerebral traumas, CC anti-iduotype antibodies.

SQ Sequence 140 AA;
PRESENTATION OF A CONTRACTOR O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ID R81330 standard; Protein; 123
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25-JAN-1995;
25-JAN-1994;
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25-JAN-1994; US-186269.
(ATHE-) ATHENA NEUROSCIENCES
Bendid MM, Jones TS, Leger
WPI; 95-269276/35.
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leukocyte adhesion to end
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Mouse anti-VLA-4 antibody 21.6 heavy chain variable region.
Humanized antibody; leukocyte adhesion molecule; VLA-4; therapeutic;
                                                                                                                                                                                                                                                                                                                                                   Key
                                                                                                                                                                                                                       region
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132. .:
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e= "framework region
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85
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Disclosure; Page 68; 105pp; English.

CC The sequence represents the mouse anti-VLA-4 antibody 21.6 heavy chain constraint framework regions of the Rel antibody for the light chain and constant framework regions of the Rel antibody for the light chain and constant framework regions of the Rel antibody for the light chain and constant framework regions of the Rel antibody for the light chain and constraint for the light chain and constant framework regions of the Rel antibody for the light chain and constraint regions. In the humanized light chain, amino acids L45, L49, constant regions. In the humanized light chain, amino acids L45, L49, constant regions. In the humanized light chain, amino acids L45, L49, constant regions. In the humanized light chain, amino acids L45, L49, constant regions. In the humanized light chain acids L45, L49, constant regions. In the humanized light chain acids L45, L49, constant regions. In the humanized light chain acids L45, L49, constant regions. In the humanized light chain acids L45, L49, constant regions. In the humanized light chain acids L45, L49, constant regions. In the humanized light chain acids L45, L49, constant regions. In the humanized light chain acids L45, L49, constant regions. In the humanized lation of a leukocyte constant regions. In the humanized lead in the treatment of stroke, constant regions are not provided acids and lation or for generating constant regions and lation or for generating constant regions.
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27-JUL-1995.
25-JAN-1995; U
25-JAN-1994; U
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WPI; 95-269276/35.
New humanised antibodies against VLA-4 - used for inhibiting
leukocyte adhesion to endothelial cells, partic. for treating
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Human VLA-4 reshaped antibody 21.6 light heavy variable region.
Humanized antibody; leukocyte adhesion molecule; VLA-4; therapeutic;
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region
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                                                                                                                                                                                                                                                    . 49
                                                              .68
e= "framework region
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                                                                                                                                                    "complementarity determining region
                                                                                                                                                                                                                  "framework region 1"
                                                                                                                                                                                                                                                                             "signal peptide'
                               "complementarity determining region
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The sequence represents the human reshaped antibody 21.6 heavy chain variable region against leukocyte adhesion molecule VLA-4. Cloned cDNA sequences of mouse 21.6 VH (099892) and vI (099889) regions are linked to human constant regions in the construction of a humanized antibody against VLA-4. The 5 and 3 ends of the mouse cDNAs are modified using PCR primers (099895-98) and then subcloned into mammalian cell expression vectors containing human subcloned into mammalian regions. In the humanized heavy chain, kappa or gamma-1 constant regions. In the humanized heavy chain, amino acids H27, H28, H29, H30, H44 and H71 in the human HC VR framework are replaced by the amino acid present in the equivalent position of the mouse 21.6 Ig H chain. Plasmids encoding the chimeric antibodies are transfected into COS cells. The humanized antibodies can be used to inhibit adhesion of a leukocyte to an endothelial cell and to treat inflammatory diseases such as multiple sclerosis. They can also be used in the treatment of stroke, also be used for detecting VLA-4, for affinity purification or for generating anti-1dioryce antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-JUL-1995
25-JAN-1995; U01219
25-JAN-1994; US-186269
(ATHE-) ATHENA NEUROSCIENCES I
Bendig MM, Jones TS, Leger O
WPI; 95-260276/35.
N-PSDB; Q99894
                                                                                                                                                                                                                                                                                                                                                                                         leukocyte adhesion to endo inflammatory disease. Disclosure; Fig 11; 105pp;
generating anti-idiotype Sequence 142 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                   New humanised antibodies against VLA-4 - used for inhibiting leukocyte adhesion to endothelial cells, partic. for treating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         W09519790-A1
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132. .1
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e= "complementarity determining region
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                         antibodies.
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R81333 Length: 142 February 11, 2000 15:48 Type: P Check: 9019

ь MDWTWRVFCL LAVAPGAHSQ VQLVQSGAEV KKPGASVKVS CKASGENIKD

GORLEWMGRI DPANGYTKYD PKFQGRVTIT ADTSASTAYM

101

ELSSLRSEDT AVYYCAREGY YGNYGVYAMD

YWGQGTLVTV

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!!AA_SEQUENCE 1.0
NA encoding tyrosine phosphatase MPTP delta - useful for DNA encoding tyrosine phosphatase MPTP delta - useful for elucidation of signal transmission mechanisms.

Claim 2; Page 11-12; 14pp; Japanese.

The sequences given in R75202-03 are encoded by insertion sequences which were included in the tyrosine phosphatase MPTP-delta coding sequence. The INS-1 CDNA was inserted between the 65th A and the 67th G of the MPTP-delta coding sequence, and the INS-2 CDNA was inserted between the 1194th A and the 1195th T of the sequence. The MPTP-delta sequence was isolated from murine brain tissue and was cloned, for expression, into the downstream region of a glutathione-S-transferase sequence and expressed as a fusion protein in E. coli. MPTP-delta
                                                                                                                                                                                                                                                                                                                                                                                                               10-MAY-1996 (first entry)

Tyrosine phosphatase MPTP-delta insertion sequence, INS-1.

Tyrosine phosphatase MPTP-delta; murine; brain tissue;

Tyrosine phosphatase MPTP-delta; murine; brain tissue;

glutathione-S-transferase; fusion protein; E. coli; differentiation; activation; information transmission; nervous system; immune system; carcinogenesis; insetion; INS-1; INS-2.

Mus musculus domesticus.

107.236487-A.
                                                                                                                                                                                                                                                                                           28-FEB-1994; 054726.
28-FEB-1994; JP-054726.
(TOKS-) TOKYOTO SHINKEI KAGAKU SOGO KENKYUSHO
WPI; 95-347455/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     standard; Protein;
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proteins regulate differentiation and activation of cells. This sequence can be used in the elucidation of the molecular mechanism for information transmission in cells, regulation mechanisms in the nervous system or immune system, or in the mechanism of carcinogenesis. Sequence 225 AA;

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R75202
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14-AUG-1986; US-896611.
12-FEB-1988; US-155245.
14-FEB-1989; US-310603.
14-FEB-1991; US-642219.
17-AUG-1993; US-108763.
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Eimeria maxima
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protection of newborn chicks from Eimeria spp. infection by immunising a laying hen prior to the hen laying with an Eimeria spp. immunising a laying hen prior to the hen laying with an Eimeria spp. gametocyte antigenic protech Example 9; Fig 14a: 39pp; English.

Example 9; Fig 14a: 39pp; English.

Example 9; Fig 14a: 39pp; English.

Example 9: Fig 14a: 3pp; English.

Exampl
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Mencher D, Pugatsc
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region
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Nsk2 receptor intracellular domain
Nsk2; neural fold/somite kinase; r
vector; tumour; muscular dystrophy
                                                                                                                                                                                                                                    R94989;
                                         region
                                                                                 region
                                                                                                                               SuM
                                                                                                                                                                                                                                                   394989 standard; Peptide; 354
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                                                                                                                             musculus.
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                                            /label- Juxtamembrane_region 60. .341
       /label= Tyrosine-kinase_region
157. .177
                                                                                                         Location/Qualifiers
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dystrophy;
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                                                                                                                                                      in (aa518-871).
receptor tyrosine kinase;
hy; diagnosis; therapy.
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Reith A;

Well-16-221725/22.

Reith A;

Well-16-221725/22.

Recording muscle receptor tyrosine kinase(s) - and pull-standard process. Transformed cells, polypeptide(s) and antibodies, related vectors, transformed cells, polypeptide(s) and antibodies, related vectors, transformed cells, polypeptide(s) and antibodies.

The tyrosine kinase region (R94990) and other polypeptide portions the tyrosine kinase region (R94990) and other polypeptide portions the tyrosine kinase region (R94990) and other polypeptide portions the tyrosine for Nsk2 in growth and differentiation, in used to study the role of Nsk2 in growth and differentiation, in the development of Nsk (ant)agonists, to identify other components the development of Nsk (ant)agonists, to identify other components tumours and muscular dystrophy. Nsk2 is a novel member of a new terminally differentiated skeletal myotubes in vitro and in adult cerminally differentiated skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in adult considered the skeletal myotubes in vitro and in the skeletal myotubes in vitro a
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25-APR-1996; U13490:
10-OCT-1994; GB-020389.
10-OCT-1994; GB-010574.
24-MAY-1995; GB-010574.
(LUDW-) LUDWIG INST CANCI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nucleic acid encoding muscle receptor tyrosine kinase(s) and related vectors, transformed cells, polypeptide(s) and antibodies, useful for treating tumours, diagnosing muscular dystrophy, etc. claim 8; Fig 2; 65pp; English.

Claim 8; Fig 2; 65pp; English.

The intracellular domain (R94989) and other polypeptide portions rate intracellular domain (R94989) and other polypeptide portions rate intracellular domain (R94989) and other polypeptide portions the development of Nsk 2 in growth and differentiation, in used to study the role of Nsk2 in growth and differentiation, in used to study the role of Nsk2 in growth and differentiation, in used to study the role of Nsk2 in growth and therepy of e.g. tumours and muscular dystrophy. Nsk2 is a novel member of a new family of muscle receptor tyrosine kinases that are expressed in terminally differentiated skeletal myotubes in vitro and in adult skeletal muscles in vivo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-JUL-1996 (first entry)
Nsk2 receptor tyrosine kinase region (aas77-858).
Nsk2 receptor tyrosine kinase;
Nsk2; neural fold/somite kinase; receptor tyrosine kinase;
vector; tumour; muscular dystrophy; diagnosis; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                            10-OCT-1995; U13490.
10-OCT-1994; GB-020389.
24-MAY-1995; GB-010574.
(LUDW-) LUDWIG INST CAN
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WO9611664-A2.
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342. .354
/label= C
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SQ R94990

Sequence

282

AA;

Length: 282 February 11, 2000 15:48

101

51

IEYVRDIGEG

REAALMAEFD NPNIVKLLGV CAVGKPMCLL

AFGRVFQARA PGLLPYEPFT MVAVKMLKEE ASADMQADFQ

Type: P

Check: 5928

TRARVSSPGP PPLSCAEQLC

IARQVAAGMA YYKADGNDAI

FEYMAYGDLN

151

DLATRNCLVG

ETMVVKIADE

GLSRNIYSAD

PIRWMPPESI YLSERKFVHR EFLRSMSPHT

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R89896
                                                                                                                                                                                                                                                                                      sequences encoding Kynurenine aminotransferase (KAT) can be inserted into vectors and subsequently cells and hence can be used for gene therapy. The vector and host cells can be used for cerebral implantation to where KAT can directly catalyse the production of kynurenic acid (KYNA) from kynurenine (KNN). It is thought KYNA acts function. KYNA concentrations and the activity of KAT show an increase with age. KAT inhibitors, by providing an increase of the glutamatergic tone at the NNDA receptor, could be useful in situations where NNDA receptor function is insufficient and/or KAT solventity and KYNA levels are abnormally enhanced. Hence they could consequences associated with the aging processes in the brain. Sequence 423 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23-JUN-1995; U07855.
07-JUL-1994; US-271667.
(PHAA ) PHARMACIA SPA
(UYMA-) UNIV MARYLAND BALTIMORE.
Benatti L. Breton J. Mosca M. Okuno E,
Speciale C:
WPI; 96-097623/10.
N-PSDB; T11560.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Isolated DNA encoding mammalian kynurenine amino:transferase (KAT) -
useful in gene therapy applications and for identifying KAT in brain
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14-701-1996 (first entry)
Rat kynurenine aminotransferase (KAT) clone.
Kynurenine aminotransferase; KAT; kynurenic acid; KYNA; kynurenine;
KYN; brain; NMDA receptor; glutamatergic function.
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                                                                                                                                                                                                                                                      Length: 423 February 11, 2000 15:48
FVKDKATLQA MDERLRKWKE
                                MPDLPGAEDE
                                                         QPSSYFLQLP
                                                                                      ATGWKVGWVM
                                                                                                                                                                        YGALFTAFQA LVDEGDEVII
                                                                                                                                                                                                                         MTKRLQARRL DGIDQNLWVE FGKLTKEYDV VNLGQGFPDF SPPDFATQAF
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                     PYDRRFAKWM IKNMGLVGIP VSTFFSRPHQ KDFDHYIRFC
                                                                                                                                          QLDPAELASK FTPRTKVLVL
                                                           QAMELNRDHM
                                                                                  GPDNIMKHLR TVHQNSIFHC
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                                                                                                             YDGHQHVSIA
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                                                    LWISQGSYFL
                                                                               PTQAQAAVAQ
                                                                                                                                                                                              GKLLGQEMDP LTNVLVTVGA
                                                                                                           SLPGMWDRTL
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                                                                             CFEREQQHFG
                                                                                                          TIGSAGKSFS
                                                                                                                                      FSRMELELVA
                                                                                                                                                                   FVTLKPSPAP
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Claim 16; Figure 3; 51pp; English.

C Sequences encoding Kynnrenine aminotransferase (KAT) can be inserted into vectors and subsequently cells and hence can be used for gene therapy. The vector and host cells can be used for cerebral implantation to where KAT can directly catalyse the production of Kynurenic acid (KYNA) from kynurenine (KYN). It is thought KYNA acts in regative endogenous modulator of cerebral glutamatergic function. KYNA concentrations and the activity of KAT show an increase with age. KAT inhibitors, by providing an increase of the concentrations where the NMDA receptor, could be useful in situations where NMDA receptor function is insufficient and/or KAT consequences associated with the treatment of the pathological consequences associated with the agine processes in the brain.

C Struct Structurally useful in the treatment of the pathological consequences associated with the agine processes in the brain.

C Sequence 437 AA;
R89897 Length: 437 February 11, 2000 15:48 Type: P Check: 206
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N-PSDB; T11742.

Isolated DNA encoding mammalian kynurenine amino:transferase (KAT) useful in gene therapy applications and for identifying KAT in brainser.
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07-JUL-1994; US-271667.
(PHAA ) PHARMACIA SPA.
(UYMA-) UNIV MARYLAND BALTIMORE.
Benatti L, Breton J, Mosca M,
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351
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SYFLIADISD FKSKMPDLPG AEDEPYDRRF
                                                                                                                                FPDFSPPDFA TQAFQQATSG NFMLNQYTRA FGYPPLTNVL
                                                                                                                                                    MNHSTWCLFF KKLTMTKRLQ ARRLDGIDQN LWVEFGKLTK EYDVVNLGQG
                                                                                           GCPVFVTLKP
                      QHFGQPSSYF
                                                                                                          TVGAYGALFT AFQALVDEGD EVIIMEPAFD CYEPMTMMAG
                                                                 ELVANLCQQH DVVCISDEVY
                                                                                        SPAPKGKLGA
                                              KSFSATGWKV
                      LQLPQAMELN
                                           GWVMGPDNIM
                                                                                        SNDWQLDPAE
                   RDHMIRSLQS
                                         KHLRTVHQNS
                                                              QWLVYDGHQH
                                                                                      LASKFTPRTK
                  VGLKLWISQG
                                      IFHCPTQAQA
                                                               VSIASLPGMW
                                                                                    VLVLNTPNNP
                                                                                                                                ASFFGKLLGQ
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25-JAN-1996.
23-JUN-1995; U07855.
07-JUL-1994; US-271667.
(PHAA ) PHARMACIA SPA.
(UYMA-) UNIV MARYLAND BALTIMORE.
                                                                Rattus rattus. WO9601893-A1.
                                                                                14-JUI-1996 (first entry)
Rat kynurenine aminotransferase (KAT) clone.
Rynurenine aminotransferase; KAT; kynurenic acid; KYNA;
KYN; brain; NMDA receptor; glutamatergic function.
                                                                                                                                                           SEQUENCE 1.0
                                                                                                                                           standard; Protein;
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                                                                                                                                           AA
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401

RPHQKDFDHY IRFCFVKDKA TLQAMDERLR KWKELQP

AKWMIKNMGL

VGIPVSTFFS

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Sequences encoding Kynurenine antinotransferase (KAT) can be inserted Sequences encoding Kynurenine antinotransferase (KAT) can be used for gene can be cectors and subsequently cells and hence can be used for gene therapy. The vector and host cells can be used for cerebral implantation to where KAT can directly catalyse the production of implantation to where KAT can directly catalyse the production of as a negative endogenous modulator of cerebral glutamatergic can concentrations and the activity of KAT show an increase with age. KAT inhibitors by providing an increase of the confirmatergic tone at the NMDA receptor, could be useful in contact tone at the NMDA receptor, could be useful in cataly where NMDA receptor function is insufficient and/or KAT consequences associated with the aping processes in the brain. Consequences associated with the aging processes in the brain. Consequences associated with the aging processes in the brain.
ALD DE PROPERTIES PROP
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                             23-NOV-1995.

18 -MAY-1995; US-245294.

RR 18-MAY-1994; US-245294.

RR 16-SEP-1994; US-307279.

R (USSH) ) US DEPT HEALTH & HUMAN SERVICES.

19 Anderson BE, Regnery RL;

10 Anderson BE, Regnery RL;

10 Nucleic acids of Rochalimaea henselae and R.quintana - methods which provided the identification of R.henselae, which is a causative agent of both cat scratch disease and bacillary angiomatosis of both cat scratch disease and bacillary angiomatosis of both cat scratch (190pp; English.

11 R Nucleic acids (190402, 1904403), fragments and antibodies of the mucleic acids (190402, 1904403), fragments and antibodies of the encoded proteins (R77433, R77434), may be used in the cangiomatosis caused by R. henselae. The proteins or fragments of them, may be used in vaccines to protect against R. henselae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Benatti L,
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W-PSDB; T11743.
Isolated DNA encoding mammalian kynurenine amino:transferase (KAT) -
useful in gene therapy applications and for identifying KAT in brain
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19-AUG-1996 (first entry)
Antigen of Rochalimaea henselae.
Rochalimaea henselae; cat scratch disease; bacillary angiomatosis;
CSD; infection; antigen; antibody; vaccine.
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TGF-a-DETA-DGAL4 multidomain protein; gene transfer; gene therapy;
nucleic acid transfer system; gene transfer; gencer;
cell targeting; multidomain protein; vector; cancer;
cell targeting; multidomain protein; vector; cancer;
exotoxin a; DETA; ompa; signal peptide; GAL4; TGF-a;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   51
                                                                                                                                                                                                                                                 W09613599-A1.
09-MAY-1996.
31-OCT-1995; F
01-NOV-1994; F
(WELS/) WELS
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        specific cell (4-65; 106pp; English. Claim 7; Page 64-65; 106pp; English. A multidommain protein (R99054) has a FLAG epitope, a portion A multidommain protein (R99054) has a FLAG epitope, a portion of human transforming growth factor-alpha (FGG-a) that acts as a of human transforming growth factorin of Pseudomonas acruginosa ligand domain, a non-cytotoxic portion of Pseudomonas acruginosa extoxin A acting as a translocation domain and the DNA exotoxin A acting as a translocation domain and the DNA exotoxin A acting as a translocation domain and the growth of a fusion binding domain of a can be expressed in E. coli (resulting in gene (729410) and can be expressed in E. coli (resulting in removal of an ompa signal peptide). It is used with an effector nucleic acid that comprises e.g. a gene to be delivered to a cell and a cognate structure for the GAL4 DNA binding domain. This provides a novel means of nucleic acid transfer, suitable for the to-rany.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            domain
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                                                                                                                                                                         Fominaya J, Wels W;
WPI; 96-239505/24.
WPSDB; T29409.
Nucleic acid transfer system for gene therapy, e.g. against cancer
- includes toxin translocation domain to target nucleic acid to
                                                                                                                                                                                                                                                                                                                                              peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pseudomonas aeruginosa;
Saccharomyces cerevisia
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Location/Qualifiers
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'label- G
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rendoplasmic
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SS

Sequence

342

AA;

Length: 342

251 201 151 101

VETDMPLTLR QHRISATSSS EESSNKGQRQ LTVSSSDYKD

standard; Protein; 421

B

CRLKKLKCSK EKPKCAKCLK NNWECRYSPK TKRSPLTRAH

REDLDMILKM DSLQDIKALL

TGLFVQDNVN

KDAVTDRLAS LTEVESRLER LSSIEQACDI GSGGDLGEAI

GWEQLEQCGY PYQRLVALYL AARLSWNQVD QVIRNALASP

DYKDDDDKLG TGVVSHFNDC PDSHTQFCFH GTCRFLVQED

February 11, 2000 15:48 Type: P

VGARCEHADL LASLEHHHHH

HLEGGSLAAL

TAHQACHLPL ETFTRHRQPR

KPACVCHSGY Check: 9286

REQPEOARLA

LTLAAAESER

FVRQGTGNDE AGAANADEKL

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WO9613599-A1.
09-MAY-1996; E04270.
31-CCT-1995; E04270.
01-NOV-1994; EP-810627.
(WELS/) WELS W;
WPI, 96-239505/24.
specific cell "Characteria commain to target nucleic acid to Claim 7; Page 67-69; 106pp; English.

A multidomain protein (R95055) has a FLAG epitope, a portion of human interleukin-2 that acts as a ligand domain, a non-cytotoxic portion of pseudomonas aeruginosa exotoxin A acting as a translocation domain and the DNA binding domain of yeast GAL4. It is the product of a fusion gene (T29411) and can be expressed in E. coli (resulting in removal of an ompA signal peptide). It is used with an effector nucleic acid that comprises e.g. a gene to be delivered to a cell and a cognate structure for the GAL4 DNA binding domain. This provides a novel means of nucleic acid transfer.
                                                                                                                          NU-PSDB; T29411.
Nucleic acid transfer system for gene therapy, e.g. against cancer
includes toxin translocation domain to target nucleic acid to
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Nucleic acid transfer system; gene transfer; gene therapy; cell targeting; multidomain protein; vector; cancer; exotoxin A; DETA; ompA; signal peptide; GAL4; interleukin:II.2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             peptide
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Chimeric Homo sapie
Chimeric Pseudomona
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Saccharomyces cerevisiae.
Location/Qualifiers
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                                                                                                                                                                                                                                                                       /note=
                                                                                                                                                                                                                                                                                                         'label- GAL4
'note- "amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'label= FLAG_epitope
                                                                                                                                                                                                                                                                                                                                    label Spacer
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note "amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                   note= "amino acids
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                                                                                                                                                                                                                                                             - Spacer
"endoplasmic reticulum retention signal"
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R91225

Length: 328

MEWDNGTGQA LGLPPTTCVY RENFKQLLLP PVYSAVLAAG

February 11, 2000 15:48

Type: P

LPLNICVITQ Check: 7811

TAVYTLNIAL ADLLYACSLP LLIYNYAQGD HWPFGDFACF

SFORYLGICH

PLAPWHKRGG

RRAAWLVCVT

FLLPFAALLA CYCLLACRLC

RQDGPAEPVA QERRGKAARM AVVVAAAFAI QRNRTVCYDL SPPALATHYM PYGMALTVIG

PTAIFAATGI HGSILFLTCI

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G-protein coupled receptor protein DNA and protein - also methods in solating (ant) agonists for treatment of cystic fibrosis, licontinence and diabetes claim 6; page 267-68; 360pp; English.

A novel human placenta-derived G-protein coupled receptor protein phate-17 (T16368). The protein can be obtd, by expression of the purinceptor in transformed host cells. It was classified as a used to screen agonists and antagonists that modulate G-pcRp sequence 328 AA;
                                                                                                                                                        WPI; 96-139698/14.
N-PSDB; T18368.
                                                                                                                                                                                                                                                                                                                                                                           26-AUG-1996 (first entry)
Human placenta G-protein coupled receptor protein.
G-protein coupled receptor protein; G-PCRP; agonist;
Cystic fibrosis; incontinence; diabetes; diagnosis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   suitable for gene therapy Sequence 421 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length: 421 February 11, 2000 15:48 Type: P Check: 7433
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                                                                                                                                                                                                                                                                                                                                                                                                                                                standard; Protein; 328 AA
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PA (UYSA-) UNIV SANKATCHEWAN.

PA (UYSA-) UNIV SANKATCHEWAN.

PI Redmond MJ. Sankos M. Campos M;

PI Redmond MJ. Sankos M;

PI Redmond M M;

P
R99695 Length: 25
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23-DEC-1992;
04-FEB-1991;
04-FEB-1992;
06-SEP-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 301 FYFTQKKFRR
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VP6 binding peptide B.
                                                                                                                                                                    27-DEC-1995; 309454.
28-DEC-1994; JP-337652.
(NITT) NITTO CHEM IND C
Mizumura Y, Yu F;
WPI; 96-302346/31.
N-PSDB; T32242.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  virus protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             VP6; binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         399695;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE 1.0
                                                                                                                                                                                                                                                                                                                                                           14-SEP-1996 (first entry)
Nitrilase regulatory factor; gene activator; promoter;
Nitrilase regulatory fator; gene activator; promoter;
Rhodococcus erythropolis strain SK92 (FERM BP-3324).
Claim 1; Page 9-10; 23pp; English.
Claim 1; Page 9-10; 23pp; English.
The nitrilase regulatory factor of Rhodococcus erythropolis SK92
The nitrilase regulatory factor of the 20pen reading frames
(R95502). These are the products of the 2 open reading frames
(R95502). These are the products of the 20pen reading frames
(R95502). These are the products of the 20pen reading frames
                                                                                                                                    enhances activation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SFLPFHITKT AYLAVGSTPG VPCTVLEAFA AAYKGTRPFA SANSVLDPIL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CNIAPASIVS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            standard; Peptide; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; Protein; 244 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     peptide; encapsulation; target; agent; drug delivery;
                                                                                                                                       erythropolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RPHELLQKLT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    February 11, 2000 15:48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RNIVYTRAQP NQDIA
                                                                                                                                       is two component nitrilase gene activator nitrilase gene promoter in presence of
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!!AA_SEQUENCE 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   of the nitrilase gene in SK92 chromosomal DNA. Introduction of a plasmid carrying the regulatory gene plus the nitrilase gene and its promoter allows efficient prodm. of nitrilase by a Rhodococus sptransformant. The nitrilase is useful for the commercial prodm. of organic acids and amides from nitriles. Sequence 244 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-MAY-1996.
21-NOV-1995, G02720.
22-NOV-1994, GB-023540.
19-JAN-1995, GB-01074.
29-JUN-1995, GB-013293.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        wasp venom BrhTX-1 subunit (b) partial sequence.
wasp; venom; neurotoxin; insecticide; biological control agent;
Lepidoptera; insect.
Bracon hebetor.
w09616171-A1.
02-UUL-1996.
23-MAR-1995; 409199.
23-MAR-1995; US-409199.
(NEWE ) NEW ENGLAND BIOLABS I
(NEWE ) X S;
X1ao J, Xu S;
WPI; 96-321120/32.
N-PSDB; T35805.
                                                                                                                                                                                                                                                                                                                                                                                        Baule VJ. Christian PD. Duncan RE. Windass JD:
Baule VJ. Christian PD. Duncan RE. Windass JD:
Baule VJ. Christian PD. Duncan RE. Windass JD:
WPSI: 96-268607/27.

WPSI: 96-268607/27.

Bracon habetor toxins and DNA encoding them - useful in biological
Bracon habetor toxins and DNA encoding them - useful in biological
Bracon habetor toxins and DNA encoding them - useful in biological
Bracon habetor toxins and DNA encoding them - useful in biological
Example 6: Fig 11: 83pp English.

Partial cDNA clone pBrhTx-1(b)1 (T32444) codes for a portion
Partial cDNA in the samplification of wasp cDNA in
BrhTx-1. It was obtd. by FCR amplification of wasp cDNA in
BrhTx-1. It was obtd. by FCR amplification of a plaque-purified
a phage clone. Primers BH(b) and BH(b)D (T32445-46) based on the clone
were used to re-screen the library beading to the isolation of clone
BrhTx-1(b) (T32429) that coded for the toxin (b) subunit (R99577).

This can be utilised in breeding of biological control agents used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        51
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Wasp venom BrhTX-1 subunit
                                                                                                                                                                                                                                                                                                                                                         Sequence
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                                                                                                          SaclM; SaclR; Streptomyces achromogenes; Streptomyces achromogenes. 055532153-A.
                                                                                                                                                                                                            w00624 standard; Protein; 358
                                                                                                                                                                                                                                                            51 GVGDCCRSHD SCGRMIKPGE TYGDVTNKGF
                                                                                                                                                                               3-0CT-1996
                                                                                                                                                                                                                                                                                           μ
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ce 83 AA;
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                                                                                                                                                                                 entry)
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                                                                                                                                                                                                                                                                      MNS
                                                                                                                                                                                                                                                                                                                                  Type: P
                                                                                                                                                    SacI; methylase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Check: 2030
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I!AA_SEQUENCE 1.0
ID R94579 standard;
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28-MAR-1996.
20-SEP-1995; J01896.
20-SEP-1995; JP-06011.
28-APR-1995; JP-106006.
28-APR-1995; JP-106006.
28-APR-1995; JP-106006.
28-APR-1995; JP-106009.
            The present sequence is the C. pneumoniae polypeptide antigen, polypeptide A. C. pneumoniae strain YK41 was cultured and genomic blaw extracted to prep. a lambda gt11 DNA library. The library was was prepd. by fusing spleen cells from a mouse infected with an anti-YK41 monoclonal antibody (MAb), which was prepd. by fusing spleen cells from a mouse infected with YK41 was both did was then fused with the expression vector pADA431 to produce a MAb expressing hydridoma. If the DNA obth was then fused with the expression vector pADA431 to host, which was cultured to give the antigenic polypeptide. The blasmid was used to transform an E coli polypeptide A. Polypeptide A and primers and probes derived from antibodies and DNA, respectively, useful in the diagnosis of semicona antibodies infection.
                                                                                                                                                                                                                                                                                                                                                                                    diagnosis
Claim 1; p
                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 96-188399/19.
N-PSDB; T14612.
Recombinant Chlamydia
                                                                                                                                                                                                                                                                                                                                                                                                                                           used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chlamydia pneumoniae polypeptide antigen (polypeptide A). Polypeptide antigen: polypeptide A; strain YK41; plasmid; pCPN533alpha; primer; assay; detection; antibody; diagnosi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chlamydia pneumoniae.
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07-NOV-1996 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA encoding SacI restriction endonuclease and methylase - useful for prodn. of the enzyme.

Claim 3; Column 15-18; 16pp; English.

This sequence represents the SacI endonuclease from Streptomyces achromogenes. W00623 represents the SacI methylase. The DNA encoding sachromogenes, and amplified using the primers shown in T35806-T35811. The amplified sequences were then cloned into the plasmid pLG339 which was used to modify an E.coli host cell. The host cell can then be used Sequence 358 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 51
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u H, Matsumoto A,
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                                                                                                                                                                                                                                                                                                                                                                                                                              detection and
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                                                                                                                                                                                                                                                                                                                                                                                                                 pneumoniae antigen and antibodies to it d assay of C. pneumoniae e.g. in clinical
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Obara
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "claimed N-terminal fragment"
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R94580

Length: 271

February 11, 2000 15:48

MSISSSSGPD

NOKNIMSOVL

UAAGKESSES

Type: P

101

AALSGKSSGS AKLETPELPK PGVTPRSEVI EIGLALAKAI QTLGEATKSA

EEASKSMEST LESLQSLSAA QMKEVEAVVV

TKTETAPQQG

P Check: 3773
IQQTRQGKNT
QKAGADTGVS

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diagnosis

rectain 2; Pages 64-66; 128pp; Japanese.

Chaim 2; Pages 64-66; 128pp; Japa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20-SEP-1994;
28-APR-1995;
28-APR-1995;
28-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                Recombinant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C. pneumoniae polypeptide antigen (polypeptide A) variant.
Polypeptide antigen; polypeptide A; strain YK41; plasmid;
pCPN533alpha; primer; assay; detection; antibody; diagnosi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            07-Nov-1996 (first entry)
C. pneumoniae polypeptide
Polypeptide antigen; polype
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                                                                                                                                                                                                                                                                                                                                                                           for detection
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AALSGKSSGS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              EMESDATIAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MSISSSGPD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; Protein; 271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pneumoniae.
                                                                                                                                                                                                                                                                                                                                                         Chlamydia pneumoniae antigen and antibodies to it tection and assay of C. pneumoniae e.g. in clinical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    JP-106008.
JP-106008.
JP-106008.
JP-106009.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  variant
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GISKVFAKGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TVQAVVQAVK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATKIAMQTSI EEASKSMEST LESLQSLSAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           February 11, 2000 15:48
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₽ G
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       diagnosis;
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LSNYASTQAQ ADQTNKLGLE KQAIKIDKER EEYQEMKAAE QKSKDLEGTM

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diagnosis

PS Example 1; Pages 75-77; 128pp; Japanese.

PS Example 1; Pages 75-77; 128pp; Japanese.

CC The present sequence is the C. pneumoniae strain YK41 was

CC polypeptide A clone, 53-3S. C. pneumoniae strain YK41 was

CC cultured and genomic DNA extracted to prep. a lambda gtil DNA

CC library. The library was then screened with an anti-YK41

CC monoclonal antibody (MAD), which was preped. by fusing spleen cells

CC from a mouse infected with YK41 with myeloma By/NS1/-B44-1 to

CC produce a MAD expressing hydridoma. The DNA obtd. was then fused

CC with the expression vector pADA431 to give pcPN533alpha. The

CC plasmid was used to transform an E. coli host, which was cultured

CC to give the antigenic polypeptide, polypeptide A. Polypeptide A

CC and primers and probes derived from its DNA can be used in assays

CC for the detection of polypeptide A antibodies and DNA,

CC respectively, useful in the diagnosis of C. pneumoniae infection.
                                                                                                                                                                                                                                                                                                                                                                                                       PRESENTATION
                                                                                                                                                                                                                                                                                                                                                                            R94586
                                                                                                                                           _SEQUENCE 1.0
R99316 standard;
R99316;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Izutsu
WPI; 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE 1.0
R94586 standard; Protein;
                                                                                                                                                                                                                                                                                       101
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C. pneumoniae polypeptide antigen (polypeptide A) clone 53-3S.
Polypeptide antigen; polypeptide A; strain YK41; plasmid; probe;
pcpN533alpha; primer; assay; detection; antibody; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    251
                                                                  13-NOV-1996 (first entry)
Human SH-PTP1 variant derived from erythroleukaemia cell
PTP; protein tyrosine phosphatase; SH2; Src homology reg-
chromosome 12p; abnormality; mutation; detection; probe;
cancer; leukaemia; diagnosis; megakaryocyte regulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 96-188399/19.
N-PSDB; T14622.
                                        US5536636-A.
                                                                                                                                                                                                  251
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          infection;
                                                                                                                                                                                                                                                                                                                                                                         Length: 259
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                                                                                                                                                                                                  DIVITYMIA
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                                                                                                                                                                                                                                                            AALSGKSSGS
                                                                                                                                                                                                                                                                                                                  EMESDAT IAG
                                                                                                                                                                                                                                                                                                                                               MSISSSSGPD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         detection
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JP-224711.
JP-106011.
JP-106006.
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721112.
US-721112.
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                                                                                                                                                                                                                                                            AKLETPELPK
                                                                                                                                                                                                                                                                                       ATKIAMQTSI
                                                                                                                                                                                                                                                                                                                 ASGKDKTSST
                                                                                                                                                                                                                                                                                                                                             NOKNIMSQVL
                                                                                                                                                        Protein;
                                                                                                                                                                                                                                                                                                                                                                       February 11, 2000 15:48
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       dia pneumoniae antigen and antibodies to and assay of C. pneumoniae e.g. in clin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Obara
                                                                                                                                                          458
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                                                                                                                                                                                                                              KQAIKIDKER EEYQEMKAAE
                                                                                                                                                                                                                                                          PGVTPRSEVI
                                                                                                                                                                                                                                                                                       EEASKSMEST LESLQSLSAA
                                                                                                                                                                                                                                                                                                                  TKTETAPQQG
                                                                                                                                                                                                                                                                                                                                             TSTPQGVPQQ
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                                                                                                                                                                                                                                                            EIGLALAKAI QTLGEATKSA
                                                                                                                                                                                                                                                                                                                  VAAGKESSES
                                                                                                                                                                                                                                                                                                                                             DKLSGNETKQ
                                                                                                                                                                                                                                                                                                                                                                         Type: P
                                                       ection; probe; neoper regulation.
                                                                                                                                                                                                                               QKSKDLEGTM
                                                                                                                                                                                                                                                                                         QMKEVEAVVV
                                                                                                                                                                                                                                                                                                                  QKAGADTGVS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       odies to it in clinical
                                                                                                                                                                                                                                                                                                                                                                            Check: 9728
                                                                                   neoplasia;
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TARRES RESERVATOR SERVICE OF THE PROPERTY OF T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         two SH2 domains) obtained from a human erythroleukaemia cell line.

The variant cDNA sequence (see T35310) lacks a sequence of two nucleotides, AG, corresponding to positions 1868 and 1869 of the wild-type sequence (see T35306). A fragment complemetary to the sequence from nucleotides 537-653 is used as a probe to detect a chromosome 12pl3 abnormality associated with neoplastic disease, in partic, acute lymphoblastic leukaemia. The probe hybridises to a part of the region coding for the two tandem SH2 domains (see R9312).

If the probe will not hybridise DNA of chromosome 12pl3 from a patient sample it is indicative of an abnormality, normally associated with neoplastic, Alternatively the wild-type SH-PFP1 or SH-PFP2 gene or protein may be used for comparison to sequenced PFP genes taken from a patient, where differences indicate an abnormality. The activity of SH-PFP1 may also be purposely altered by mutation to effect a change in megakaryocyte function and hence platelet production.
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01-DEC-1992; US-983926.
28-FEB-1994; US-983926.
(BETH-) BETH ISRAEL HOSPITAL ASSOC.
(MASI ) MASSACHUSETTS INST TECHNOLOGY.
(MASI ) MASSACHUSETTS INST TECHNOLOGY.
Freeman RM, Neel BG, Plutzky J, Rose.
WPI; 96-341506/34.
N-PSDB; T35310.
                                                  23-NOV-1995
17-MAY-1995; U06244.
18-MAY-1994; US-245356.
(CHIL-) CHILDRENS HOSPITAL M
GOHINSKI RJ. WHITSEKT JA;
WPI; 96-011078/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    401
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                                                                                                                                                                                                                                                                             27-NOV-1996 (first entry)
Human thyroid transcription factor: TIF-1; human adenocarcinoma cell line;
Thyroid transcription factor: TIF-1; human adenocarcinoma cell line;
H441; rat; mouse; pulmonary adenocarcinoma; H820; small cell carcinoma;
H345; tracheal-bronchial epithelial cell lines; respiratory epithelium;
fetal lung; gestation; pro-SP-C; respiratory epithelial cell;
nonotilated bronchiolar cell; immature lung; alveolar; Type II;
epithelial cell; nonciliated; Type I; adult.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    _SEQUENCE 1.0
R83015 stand
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 standard; Protein; 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EESEEAAVSR
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polynucleotide(s)
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                                                                                                               MEDICAL
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lung
cell
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   nuclear
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Pep-

W09627007-A1.

06-SEP-1996.

PF 22-FEB-1995; AU0094...

PR 02-MAR-1995; AU-001457.

PR 20-NOV-1995; AU-006647.

PR 22-DEC-1995; AU-007274.

'AMEA-) AMEAD OPERATIONS PTY LTD.

'AMEA-) AMEAD OPERATIONS PTY LTD.
                 PT - useful for cancer diagnosis and therapy.

Sclaim 50; Fig 39; 157pp; English.

CC This sequence represents the thyroid transcription factor, TTF-1. The CT TF-1 DNA sequence was isolated from the human adenocarcinoma cell line CC H441. The TTF-1 locus is contained within a 4.6 kb BamHI fragment and CC consists of two exons and one intron. This predicted amino acid CC sequence of human TTF-1 shows close identity with the amino acid CC sequence predicted by the rat TTF-1 cDNA sequence and 92.4% identity CC with the rat TTF-1 cDNA. The intron is approx. I kb in length and CC is flanked by consensus splice donor-acceptor sites that fit splice CC acceptor-donor rules. The mRNA produced is 2.3 kb as detected by CC adenocarcinoma cells. TTF-1 mRNA was detected in human pulmonary CC adenocarcinoma cells. TTF-1 mRNA was detected in human pulmonary CC adenocarcinoma cells. TTF-1 mRNA was detected in human pulmonary CC adenocarcinoma cells. TTF-1 has been detected in nuclei of the respiratory epithelian to the for pro-SP-C. TTF-1 was detected in subsets of conscillated bronchiolar, and rarely in noncillated bronchiolar respiratory epithelial cells in the developing lung, including calveolar Type II epithelial cells. TTF-1 was not detected in subsets of alveolar Type II epithelial cells. The detected in the adult lung, and was most prominent in type II epithelial cells but was excluded from type I cells. Or cillated bronchiolar epithelial cells. The adult lung, and was most prominent in type II epithelial cells but was excluded from the tells.
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W00725 standard; Protein; 207 AA.
                                                                                                                                                                                                                                                                                                                                                                              30-NOV-1996 (first entry)
Vascular endothelial growth factor-like protein SOM175.
Vascular endothelial growth factor; VEGF; VEGF165; SOM175; neuron;
ustroglial proliferation.
                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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growth factor related to vascular endothelial growth for inducing astroglial proliferation and promoting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length: 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAAISVGSGG
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1. .21
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                                                                                                             Nordenskjold
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  factor -
ng neuronal
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R99253 Length: 374 February 11,

101

GTVDPLTKLV LVNAIYFKGK WNEQFDRKYT RGMLFKTNEE

YKDGDIHRGF QSLLSEVNRT
FYQAELEELS FAEDTEECRK

GTQYLLRTAN HINDWVAEKT

P Check: 4145
MVFMGAKGST
RLFGEKTCDF

KKTVQMMFKE

FAISLFKILG EEDNSRNVFF SPMSISSALA

2000 15:48

Type: P

5

MDDLCEANGT AAQMSQALCL

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W00725
       Human Cytoplasmic antiproteinase-2 (CAP-2) and CAP-3 - serine protease inhibitors useful in the purification of proteins and in the treatment of inflammatory diseases and apoptosis. Claim 9: Page 37-38: 50pp: English.

Human cytoplasmic antiproteinase-2 protein (CAP-2) (R99253) is an inhibitor of specific trypsin-like serine proteases. Its amino acid sequence was deduced from a CDNA clone (T35220) isolated from a human placenta lambda gtil cDNA library. Another serpin, CAP-3 (R99254), has also been identified. The CAPs are useful for the purification of proteins and in the treatment of inflammatory diseases and apoptosis. They can be expressed in transformed or 171 and 
   Sequence
                                                                                                                                                                                                                                                                                                                                                                                                N-PSDB; T35220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             02-FEB-1996; U01288
08-FEB-1995; US-385500
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05-DEC-1996 (first entry)
Cytoplasmic antiproteinase-2 protein.
Cytoplasmic antiproteinase-2 protein; CAP-2; serpin;
serine protease inhibitor; antiinflammatory; apoptosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      modified_site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             proliferation and secuence 207 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           standard; Protein; 374 AA
374 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /label N-glycosylation_site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /label= N-glycosylation_site
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ated from
n, CAP-3
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KAWTNSEKLT KSKVQVFLPR LKLEESYDLE PFLRRLGMID AFDEAKADFS

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W01743 Length: 259
                                                                                                                                                                   Homo sapiens.

06-SBP-1996.
01-MAR-1996. U02957.
01-MAR-1995. US-397651.
01-MAR-1995. US-469427.
06-DEC-1995. US-569063.
(LUDW-) LUDWIG INST CARCER RES.
(LUDW-) LUDWIG INST CARCER SES.
(LUTHE-) UNIV HELSINKI LICENSING LTD OY.
Alitalo K, Eriksson U, Olofsson B, II
WPI, 96-412552/41.

N-PSDB. T37915.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Detection and determination of anti-Chlamydia pneumoniae antibody using the polypeptide C as the antigen Example 3; Page 17-18; 18pp; Japannese. The protein is that of the Chlamydia pneumoniae 53 kDa antigen. A method for the detection and determination of anti-C. pneumoniae antibodies in a sample comprises using at least 5 consecutive amino acids of the polypeptide C 73 kDa antigen (W01742). Sequence 259 AA;
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28-APR-1995; JP-106014.
(HITB) HITACHI CHEM CO LTD
WPI; 97-056178/06.
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28-APR-1997 (first entry)
Vascular endothelial growth factor. B186.
Vascular endothelial growth factor; VPF.
Endothelial cell; proliferation; vascular endothelial growth factor; verscular endothelium; mesodermal cell; cationic dimer; tissue regeneration;
VEGF; endothelium; mesodermal cell mitogen; anglogenesis; cell growth;
vascular permeability factor; cell growth; tissue reorganisation; antibody;
embryonic development; wound healing; tissue reorganisation; antibody;
cancer; metastatic risk; tumour cell; mouse.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE 1.0
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vascular endothelial growth factor VEGF-B proteins - useful to vascular endothelial growth factor VEGF-B proteins - useful to accelerate angiogenesis in wound healing, also related nucleic acid accelerate angiogenesis in wound healing, also related nucleic acid and intibodies for cancer diagnosts.

Claim 18; Page 62; 107pp; English.

Claim 18; Page 62; 107pp; English.

WO4824-W04831 represent the vascular endothelial growth factor (VEGF) proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins of the invention, which promote endothelial or mesodermal cell proteins.
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e 53 kDa antigen.
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3, Pajusola K;
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                                                                                                                                         proteins - useful to also related nucleic acid
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IIAA_SEQUENCE 1.0

WI1481;
AC WI1480-85) Car
Sequence 200;
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답중답
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D. immitis mature venom allergen antigen
Venom allergen antigen 5-like gene; VA5;
DDIVA615; PDIVA205; vaccine; heartworm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             151
                                                                                                                                                                                                                                                                                                                                          Nucleic acid encoding helminth venom allergen antigen 5-like protein Nucleic acid encoding helminth venom allergen antigen 5-like protein pref. from Dirofilaria immitis or Onchocerca volvulus, useful esp. in vaccines to prevent helminth infection in vaccines to prevent helminth infection. Claim 19: Page 91-92; 124pp; English.

Claim 19: Page 91-92; 124pp; English.

Dirofilaria immitis mature venom allergen antigen 5-like protein va5 (W11481), or PDIVA205, is capable of eliciting an immune response (cellular and/or humoral) in animals. It is encoded by a response (cellular and/or humoral) in animals. It is encoded by a CDNA molecule nDIVA615 (751379). VA5-like proteins (see also CDNA molecule nDIVA615 (751379).
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23-MAY-1995; US-450
                                 SEQUENCE 1.0
W11479 standard; Protein; 221
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                                                                                                  GLCKK
                                                                                                                                                                                                                         WDCKLEKSAQ
                                                                                                                                                                                                                                                                  YECEGGKLTP
                                                                                                                                         DHNIVMVART
                                                                                                                                                                                  TDAGRLWWSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -1995; US-450944.
) HESKA CORP.
CA, Wisnewski N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; Protein;
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                                                                                                                                                                                                                                                                  TERKNIVTQI NKYRSRLIRG
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                                                                                                                                              LVFICHYFPG
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                                                                                                                                                GNMVKDLIYE
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                                                                                                                                                                                                                                SPSSERRGIG
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                                                                                                                                                                                         NILHFTQMAW GETYKLGSGV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5-like protein PDiVA205 helminth; parasite;
                                                                                                                                                                                                                                                                                                                  Type: P
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                                                                                                                                                                                                                                       SVRDLKKTAG
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W11479; 22-APR-1997 (first entry

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OI A WWW. SOR SERVER SE
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microorganism producing nitrilase in absence of inducer - useful as reatalyst in large scale organic acid production

claim 7; Pages 7-8; 13pp; Japanese.

The present sequence is one of the two components of the resent sequence is one of the two components of the resent sequence is one of the two components of the reconstanting the resent sequence are regulatory factor (RF), which containing the RF gene can be used to transform a microorganism, or containing the RF gene can be used to transform a microorganism, or can builting it to produce nitrilase in pright yield without adding any conductor to the medium. Hence, the preparation of the medium is easy confidered to the rest culture of the microorganism, and the cidsposal cost of the waste culture is reduced. Nutrilase is caids, as it hydrolyses nitriles into their corresponding acids.

Sequence 244 AA;
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N-PSDB, T51376;

N-PSDB, T51378.

N-PSDB, T51378.

Theref. from Dirofilaria immittle or onchocerca volvulus, useful esp. or o
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19-JUL-1995;
19-JUL-1995;
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02-JUN-1997 (first entry)
Rhodococcus erythropolis SK92-Bl regulatory factor component.
Regulatory factor; activation; nitrilase; gene promoter; recombinant plasmid; transformation; microorganism; catalyst; production; organic acid.
Rhodococcus erythropolis.
J09028382-A.
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W13272 standa
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28-NOV-1996
23-MAY-1996; U07709.
23-MAY-1995; US-450944.
(HESK-) HESKA CORP.
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Venom allergen an
nDiVA833; PDiVA2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      JUL-1995; 204061.
JUL-1995; JP-204061.
JUL-1995; JP-204061.
JUL-1995; JP-204061.
JUL-1995; JP-159092/15.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard; Protein; 244
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ergen antigen 5-like gene; vAS;
PDIVA221; vaccine; heartworm.
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/note "mature prot
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17. .221
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PN ACC DT DT DE KW
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                                                                                                                                                                                                                                                                          pT Helicobacter Pylori CagI polynucleotide and related proteins - used pT in diagnosis and in vaccines for the treatment of H. pylori infection associated disease PS Disclosure; Fig 41; 303pp; English.

CC The present sequence is encoded by a putative open reading frame (ORF) of CC putative ORFs with different polarities. It is hypothesised that some of CC putative ORFs with motification polarities with homology to the ptl genes of Bordetella pertussis and VIR B4 genes of Agrobacterium tumefactens and CC Salmonella genus. The absence of the CagA gene in the type I strains is factors restricted to type I strains). The CagI nucleotide sequence, its fargments and encoded proteins are used in the diagnosis of H. pylori type I strain) infection in an individual and in vaccines duodenal and gastric ulcers, severe forms of active gastriitis (esp. type Sequence 140 AA;
                                                                                                                                                                                                                                                        W06943
                      JO-JUN 1997 (first entry)
CagI locus product 6.
CagI; CagA; virulence factor; ej
Bordetella pertussis; VIR B4; Ac
Salmonella; type I strain; viru
vaccine; treatment; duodenal; g
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adenocarcinoma.
Helicobacter pylori.
WO9633274-A1.
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20-APR-1995; US-425194.
07-JUN-1995; US-477451.
(BIOC-) BIOCINE SPA.
COVACCI A;
                                                                                                                      v06935;
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CagI locus product 14.

CagI; CagA; virulence factor; exporter molecule; homology; ptl

Bordetella pertussis; VIR 84; Agrobacterium tumefaciens; invas

Salmonella; type I strain; virulence; diagnosis; H. pytori; in

vaccine; treatment; duodenal; gastric ulcer; active gastritis;
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N-PSDB; T46159.
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                                                                                                                                standard;
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                                                                                                                                                                      MGDYGPNNPF
                                                                                                                                                                                               LEQKVAILND
                                                                                                                                                                                                                        LLGVLSLACL
                                                                                                                           Protein;
                                                                                                                                                                                                                                               February 11, 2000 15:48
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                                                                                                                             136
                         or; exporter molecule; homology; ptl
B4; Agrobacterium tumefaciens; invassi
virulence; diagnosis; H. pylori; inf
al; gastric ulcer; active gastritis;
                                                                                                                                                                   MYGYAPTYSD
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invasion factor;
i; infection;
                                      ptl gene;
vasion factor;
infection;
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pr Helicobacter pylori CagI polynucleotide and related proteins - used pri in diagnosis and in vaccines for the treatment of H. pylori pri in diagnosis and in vaccines for the treatment of H. pylori pri indiagnosis and in vaccines for the treatment of H. pylori pri indiagnosis and in vaccines for the CagI region contains clusters of Sisclosure; Fig 4D; 303pp; English.

Provided the CagI locus (see T46159). The CagI region contains clusters of the CagI locus (see T46159). The CagI region contains clusters of CC putative ONFS with different polarities. It is hypothesised that some of CC putative ONFS with different polarities. It is hypothesised that some of CC putative ONFS with motifs shared by the purported invasion factors of CC associated with the absence of the CagA gene in the type I strains is Salmonella genus. The absence of CagI sequences (which may encode virulence associated with the absence of CagI sequences (which may encode virulence CC (esp. H. pylori type I strains). The CagI nucleotide sequence, its fragments and encoded proteins are used in the diagnosis of H. pylori cycle I strains) infection in an individual and in vaccines (esp. H. pylori type I strain) infection in factive gastritis (esp. type CC daudeinal and gastric ulcers, severe forms of active gastritis (esp. type CC castritis) and dastric adenocarcinoma.
      20-APR-1995; US-425194.

R 07-JUN-1995; US-477451.

R 07-JUN-1995; US-477451.

COVACC1 A:

WPI; 96-46--
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pr Helicobacter pylori nucleic acid sequences and related pr polypeptide(s) - useful for vaccines to treat or prevent H. pylori pr polypeptide(s) - useful for vaccines to treat or prevent H. pylori printection, and to detect Helicobacter printection, and to detect Helicobacter printection, and to detect Helicobacter contain 72; page 725-726; 1481pp; English.

CC Infection or to identify H. pylori polypeptide binding compounds, confection or to identify H. pylori polypeptide binding compounds, consecuted as potential H. pylori life cycle activators or inhibitors.

CC Unsetil as potential H. pylori life cycle activators or inhibitors.

CC overlapping contides generated by mechanically shearing the bacterial coverlapping contides generated by mechanically shearing the bacterial coverlapping contides were analysed for ORF of at least 180 nucleotides, on the sequences were analysed for oRF of at least 180 nucleotides, conditions the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the amino conditions of the pylori antigens for vaccine development, the pylori antigens for vaccine development, the pylori antigens for vaccine development, 
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N-PSDB; T46159.
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06-JUN-1996; US-487032.
07-JUN-1995; US-487032.
01-APR-1996; US-630405.
01-APR-1996; US-630405.
ASTR AB:
BEGGIANDA AB:
BEGGIANDA OF SMITH D,
WPI: 97-052306/05.
N-PSDB; T67714.
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gastritis) and gastric adenocarcinoma.
Sequence 136 AA;
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H. pylori secreted or periplasmic protein 80257.aa.
Cytoplasmic; vaccine; prevention; treatment; infection; identification; cytoplasmic; vaccine; prevention; life cycle; activator; bacteria; inhibitor; binding compound; bacterium; life cycle; activator; bacteria; diadnosis; envelope.
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diagnosis.

bacteria; inhibitor;

Helicobacter pylori. w09640893-A1.

19-DEC-1996. 06-JUN-1996; U09122. 07-JUN-1995; US-487032. 01-APR-1996; US-630405.

!!AA_SEQUENCE 1.0

standard; Protein;

276

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W20776;

15-UL-1997 (first entry)
H. pylori flagella-associated protein, 07ge20415orf34.
H. pylori flagella-associated protein; infection; envelope: Cytoplasmic; vaccine; prevention; treatment; life cycle; activato identification; binding compound; bacterium; life cycle; activato identification; binding compound; bacteria; inhibitor; duodenal ulcer disease; chronic gastritis; bacteria; inhibitor; duodenal ulcer disease; chronic

101 151

SFQPNDNLGI

NAMWGIQNLL MSQMMSNYGP NNSFMYGYAP TYSDSSFLPP

51

VSRDQLKIEI KSMKTNFYKI February 11,

PKETLEQKVA KLLFAWCLII

ILNDYNDKNV NIKFDDISLG

GMFNAPLNAD

W20886

Length:

154

2000 15:48

Type: P

Check: 2872

NNAKNKNKGV

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ID W20886 standa
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DR N.PSDB; T68139.

In Helicobacter pylori nucleic acid sequences and related prolypeptide(s) useful for vaccines to treat or prevent H. pylori prolypeptide(s) useful for vaccines to treat or prevent H. pylori prolypeptide pylori nifection, and to detect Helicobacter for claim 72; Page 1289; 1481.pp; English.

CC Infaction or to 1289; 1481.pp; English.

CC Infection or to identify H. pylori secreted or periplasmic protein.

CC useful as potential H. pylori polypeptide binding compounds.

CC useful as potential H. pylori | NTCC 5579) was determined from crown or to proper analysed for ORT of at least 180 nucleotides.

CC DNA. The sequences were analysed for ORT of at least 180 nucleotides.

CC and the predicted coding regions defined by computer evaluation. To compute development, the amino crown of the pylori antigens for vaccine development, the amino crown sequences predicted from various ORT were analysed for significant acid sequences predicted from various ORT were analysed for significant conditions the sequences of interest, particular regions can be solared from H. pylori by PCR amplification for recombinant polypeptide correduction, e.g. in E. coli hosts.
                                                                                                                                                                                                                                                                                                                                     weI; 97-052306/05.
N-PSDB; T68139.
Helicobactor
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07-JUN-1995; US-487032.
01-APR-1996; US-630405.
(ASTR ) ASTRA AB.
(ASTR ) ASTRA AB.
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Cytoplasmic; vaccine; prevention; treatment; infection; envelope; identification; binding compound; bacterium; life cycle; activator; bacteria; inhibitor; duodenal ulcer disease; chronic gastritis;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FLFALIIKPS FFYYTTYLLL LVSLSIISKY YLLSHANFTM
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PT Helicobacter pylori nucleic acid sequences and related problypeptide(s) - useful for vaccines to treat or prevent H. pylori PT infection, and to detect Helicobacter Claim 72; Page 463; 1481pp; English.

CC This sequence is a H. pylori secreted or periplasmic protein may be used in a vaccine to prevent or treat H. pylori CC infection or to identify H. pylori polypeptide binding compounds, curve the protein may be used in a vaccine to prevent or treat H. pylori CC useful as potential H. pylori life cycle activators or inhibitors.

CC overlapping contigs generated by mechanically shearing the bacterial CC useful as potenties were analysed for ORF of at least 180 nucleotides, created the predicted coding regions defined by computer evaluation. To CC and the predicted coding regions defined by computer evaluation. To CC identify likely H. pylori antigens for vaccine development, the amino CC considered protected from various ORF were analysed for significant CC and determined the sequences of interest, particular regions can be considered from H. pylori by PCR amplification for recombinant polypeptide cc isolated from H. pylori by PCR amplification for recombinant polypeptide
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CC This sequence represents a H. pylori flagella-associated protein. CC The protein may be used in a vaccine to prevent or treat H. pylori CC Useful as potential H. pylori polypeptide binding compounds, the genomic sequence of H. pylori (ATCC 5579) was determined from CC Owal as potential H. pylori (ATCC 5579) was determined from CC DNA. The sequences were analysed for ORF of at least 180 nucleotides, and the predicted coding regions defined by computer evaluation. To contify likely H. pylori antigens defined by computer evaluation. To acid sequences predicted from various ORF were analysed for significant and determined the sequences of interest, particular regions can be production, e.g. in E. coli hosts.
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07-JUN-1995; US-48
01-APR-1996; US-63
(ASTR) ASTRA AB
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WPI; 97-052306/05.
N-PSDB; T67473.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  H. pylori secreted or periplasmic protein 23564012.aa. Cytoplasmic; vaccine; prevention; treatment; infection; identification; binding compound; bacterium; life cycle; activator; bacteria; inhibitor; duodenal ulcer disease; chronic gastritis; diagnosis; envelope.
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Berglindh OT, Smit
WPI; 97-052306/05.
N-PSDB; T68029.
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; US-487032.
; US-630405.
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Smith
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PT Helicobacter Pylori nucleic acid sequences and related PT polypeptide(s) - useful for vaccines to treat or prevent H. pylori PT infection, and to detect Helicobacter PS claim 56; Pages 516; 1481pp; English.

CC The present sequence is a Helicobacter pylori cell envelope protein. CC The present sequence is a Helicobacter pylori cell envelope protein. CC infection or to identify H. pylori polypeptide binding compounds, useful cas potential H. pylori life cycle activators or inhibitors. The genomic contigs generated by mechanically shearing the bacterial DNA. The ccontigs generated by mechanically shearing the bacterial DNA. The predicted coding regions defined by computer evaluation. To identify compounds predicted from various ORF were analysed for sequences predicted from various ORF were analysed for significant compounds the sequences of interest, particular regions can be production, e.g. in E. coli hosts.
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WPI; 97-05
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-UUI-1997 (first entry)
H. Pylori cell envelope transporter protein, 2461062.aa.
Cytoplasmic; vaccine; prevention; treatment; infection; identification;
binding compound; bacterium; life cycle; activator; bacteria; inhibitor
duodenal lucer disease; chronic gastritis; diagnosis; envelope.
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07-JUN-1995;
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WO9640893-A1.
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W20324 standard; protein; 102 AA.
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Length: 102 February 11,
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97-052306/05.
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                                                                                                                                                                                                                                                                                                                                                                                                   ASTRA AB
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; US-630405.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Check: 5141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      inhibitor;
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Helicobacter pylori.
W09640893 Al
19-DEC-1996
06-DUN-1996; U09122
07-JUN-1995; US-487032
01-APR-1996; US-630405
                                                               W20090;
15-JUL-1997 (first entry)
H. pylori secreted or periplasmic protein 05ae20220orf50.
Secretion; vaccine; prevention; treatment; infection; identification; binding compound; bacterium; life cycle; activator; bacteria; inhibit
                                                                                                                                                                         SEQUENCE 1.0
                                                                                                                                                                                                          101
                                                                                                                                                                                                          Z
                                                                                                                                                                                                                     EIAEKFKLIK DKALVTVINK GYGKNPLTKN YNIKNYGELE RVIKKLPLVR
                                                                                                                                                                                                                                            MMAHSLILVS KTSLSNLLIF VVQPDGKLSM TDAAIDPNMT NSGLRWYRVN
                                                                                                                                                                standard; Protein; 121 AA.
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inhibitor;

2000 15:48 Type: P Check: 6372

W20696

Length: 121

101

SHANFTMKLI IALFSYGAGV

51

SEQUENCE 1.0

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pr polypeptide(s) - useful for vaccines to treat or prevent H. pylori prolypeptide(s) - useful for vaccines to treat or prevent H. pylori prinfection, and to detect Helicobacter

pr infection, and to detect Helicobacter

pr claim 72 Page 1115; 1481pp; English.

CC The present sequence represents a Helicobacter pylori protein which

CC is likely to be secreted or periplasmic.

CC infection or to identify H. pylori polypeptide binding compounds,

CC useful as potential H. pylori life cycle activators or inhibitors.

CC useful as potential H. pylori life cycle activators or inhibitors.

CC overlapping contigs generated by mechanically shearing the bacterial

CC overlapping contigs generated by mechanically shearing the bacterial

CC overlapping contigs generated by mechanically shearing the bacterial

CC overlapping contigs generated by mechanically shearing the bacterial

CC overlapping contigs generated by mechanically shearing the bacterial

CC overlapping contigs generated by mechanically shearing the bacterial

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CC overlapping contigs generated by mechanically shearing the bacterial

CC overlapping contigs generated by mechanically shearing the bacterial

CC overlapping contigs generated by mechanically shearing the bacter
                          Prelicobacter pylori nucleic acid sequences and related problypetide(s) - useful for vaccines to treat or prevent H. pylori probaptide(s) - useful for vaccines to treat or prevent H. pylori probaptide(s) - useful for vaccines to treat or prevent H. pylori protein fine protein for the present sequence is a H. pylori cytoplasmic protein involved in the protein may be used in a vaccine to prevent or treat H. pylori carbohydrate metabolism.

CC carbohydrate metabolism.

CC infection or to identify H. pylori polypeptide binding compounds, in the control of the pylori (AFCC 55679) was determined from the genomic sequence of H. pylori (AFCC 55679) was determined from the genomic sequences of H. pylori (AFCC 55679) was determined from the pylori contigs generated by mechanically shearing the bacterial control of the predicted coding regions defined by computer evaluation. To call the predicted coding regions defined by computer evaluation. To call sequences predicted from various of were analysed for significant acid sequences predicted from various of were analysed for significant condigy to other known or exported membrane proteins. Having identified sequences of interest, particular regions can be considered from H. pylori by PCR amplification for recombinant polypeptide sequence 53 AA;

Sequence 53 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      06-JUN-1996; U09122.
07-JUN-1995; US-487032.
01-APR-1996; US-630405.
(ASTR ) ASTRA AB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (ASTR ) ASTRA AB.
Berglindh OT, Smith D,
WPI; 97-052306/05.
N-PSDB; T67382.
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H. pylori cytoplasmic protein, 14257751.aa.

Gytoplasmic; vaccine; prevention; treatment; infection; identification; cytoplasmic; vaccine; prevention; treatment; infection; bacteria, inhibitor; binding compound; bacterium; life cycle; activator; bacteria, inhibitor; diagnosis; envelope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Helicobacter pylori.
WO9640893-Al.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TSSFKTKMKR PISKLKQNFL QFKHSFNKHL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ILVYPILELE ALIIKPSEFY YTTYLLLLVS LSIISKYYLL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   wew isolated human placenta purinergic P-2u receptor (PNR) gene used to develop prods. for the diagnosis and treatment of conditions proceed and the second expression of the receptor e.g. inflammation PS Claim 1; Page 27-29; 36pp; English.

CC W09433 shows a human placenta-derived purinergic P-2u receptor community. An assay for up-regulated expression of PNR can accelerate diagnosis and proper treatment of conditions caused by abnormal signal CC dransduction due to systemic and local infections, traumatic and cother tissue damage, hereditary or environmental diseases associated cor pathological problems.

Sequence 328 AA;
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ID W09433 standard;
W09433
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Homo sapiens.

MO9638558-A2.
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02-JUN-1995; US-459046.
(INCY-) INCYTE PHARM INC.
AU-YOUNG J, COLEMAN R,
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                                               US5635384-A.
03-JUN-1997.
11-JUN-1990;
11-JUN-1990;
109-DEC-1992;
26-JAN-1995;
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Luffin-A.
                                                                                                                                                                                                                                            Key
region
                                                                                                                                                                                                                                                                                                                pro-Ribosome Inactivating Protein; proRIP; peptide linker; cancer; pro-Ribosome; eukaryotto ribosome; alpha fragment; beta fragment; inactivation; eukaryotto ribosome; alpha fragment; beta fragment; inhibitor; protein synthesis; N-91ycosidase; 91ycosidase; flycosidase; alpha fragment; ribosomal 288 RNA; cellular proliferation; HIV-infected T cell.
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US-535636.
US-987927.
US-378761.
                                                                                                                                                                                            Location/Qualifiers
138..148
/note= "Position of F
peptide linker sequer
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                                                                                                                                                                                                  on of possible insertion sequence"
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fibrosis; hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FMVLKTILGL SHRGVRSYPM
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Length:

5 3

February

2000 15:48

Type:

Check:

Morgan AER,

Walsh

TA;

of internal

be used

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PP US5635384-A.

Peptide linker sequence

PP D3-JUN-1997.

PF 11-JUN-1990; US-35636.

PR 11-JUN-1990; US-35636.

PR 26-JAN-1995; US-378761.

PR 26-JAN-1995; US-378761.

PA (DOWC) DOWELANCO.

PI D. Morgan AER, Walsh TA;

PR 27-30931/28.

PI Inactive precursor of maize ribosome-inactivating protein - also chimeric ribosome-inactivating protein precursors containing

PT Inactive precursor of maize ribosome-inactivating protein of the chimeric ribosome-inactivating protein precursors containing

PT Inactive precursor of maize ribosome-inactivating protein precursors containing

PT Inactive precursor of maize ribosome-inactivating protein precursor sequence inactivating containing in the construction of the protein sequence in mactivating enterpolic ribosomes, but can be converted by removable, contrivating eukaryotic ribosomes, but can be converted by removal containing enterpolic protein synthesis. They possess a highly contained the sequence is a fighly contained the sequence in the construction of cells, e.g. cancer cells and HTV-infected T contains proliferation of cells, e.g. cancer cells and HTV-infected T cells. The inactive proRIP proteins make it possible to provide protein cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pri Inactive precursor of maize ribosome-inactivating protein - also prichimeric ribosome-inactivating protein precursors containing internal linker sequences
PT internal linker sequences
PS Claim 2; Column 117-118; 121pp; English.

CC The sequences given in W21698-710 represent Ribosome Inactivating CC Proteins (RIP's), which may be used in the construction of the protein for the invention. The proRIP has a selectively removable, in a converted by removal contractivating eukaryotic ribosomes, but can be converted by removal CC inactivating eukaryotic ribosomes. RIPs are potent contractivating eukaryotic ribosomes. RIPs are potent inhibitors of eukaryotic protein synthesis. RIPs are potent contractic of eukaryotic protein synthesis. RIPs are potent contractic n-q1ycosidase activity which cleaves the glycosidic bond of calenine 4324 of rat liver ribosomal 28S RNA. RIP's selectively inhibit contractivation of cells, e.g. cancer cells and HIV-infected T contraction of cells, e.g. cancer cells and HIV-infected T contraction in the synthesis inhibitors with uses in practical and improved ways not before soyible. The RIP can be used to make cytotoxic conjugates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            rat; r
Luffa
Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pro-Ribosome Inactivating Protein; proRIP; peptide linker; cancer; inactivation; eukaryotic ribosome; alpha fragment; beta fragment; inhibitor; protein synthesis; N-glycosidase; glycosidic bond; liver; rat; ribosomal 28S RNA; cellular proliferation; HIV-infected T cell.
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W21709 stand
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26-SEP-1997 (first entry)
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/note= "Position of possible insertion peptide linker sequence"
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I!AA_SEQUENCE 1.0

AID W10031;
AC W10031;
DT 24-SEP-1997 (
DE Protein encode
KW Lock-docking;
KW maize; Fusarii
OS 2ca mays.
PN W09640998-A1
19-DEC-1996;
PF 05-JUN-1995; [
PR 07-JUN-1995; [
PR 07-JUN-1996; [
PR
W maize; Fusarium moniliforme.

Sea mays; Fusarium moniliforme.

N w0940998-A1.

PD 19-DEC-1996.

PF 05-UNN-1995; UNS-481687.

PR (PION-) PIONEER HI-BRED INT INC.

PR HI1995; UNS-481687.

PR (PION-) PIONEER HI-BRED INT INC.

PR W11997-087067/08.

PR N-SDBS; 770121.

PT Method for prodn. of cDNA libraries with anchored ends - useful for subtractive cloning of sequences of interest

PS Subtractive cloning of sequences of interest

CC This protein is the longest putative open reading frame encoded by cDNA clone was identified from a full-

CC clone G-4-5 which was activated upon Fusarium moniliforme infection of close sized point cDNA clone was identified from a full-

CC designated ICEFAL (isolation of cDNA ends from an anchored library). The close sized pCR products which do not necessarily require further subcloning/

CC method of generating a cDNA library with anchored ends using a method of isolation of companies of produces discrept companies as opposed to specific cDNA clones as produced by cDNA clone was identified from a nanchored library). The companies of the produces of the produces discrept companies as opposed to specific cDNA clones as produced by cDNA clones as produced by cDNA clones full-length cDNA sobtainable from companies as opposed to specific cDNA clones, as produced by cDNA clones full-length cDNA sobtainable from companies as opposed to specific cDNA clones, as produced by cDNA clones as pCR and RACE require accompanies and companies as pcroduced by cDNA clones full-length cDNA as pcroduced by cDNA clones full-length cDNA sobtainable from companies as opposed to specific cDNA clones, as produced by cDNA clones as pcroduced by cDNA clones and RACE require as pcroduced by cDNA clones as pcroduced by cDNA clones as p
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သည်
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24-SEP-1997 (first entry)
Protein encoded by clone G-4-5 from
primer; polymerase chain reaction; c
lock-docking; screening; PCR based c
maize; Fusarium moniliforme.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               151
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Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FLVILQTTAE
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250 AA;
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cDNA library; anchored end; pcsyB;
cDNA subtractive cloning; Zea mays;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PSPAALSLEN
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IIAA_SEQUENCE 1.0

ID W13647 standard; peptide; 30 AA.

AC W13647;

DT 25-NOV-1997 (first entry)

DE Invariant Chain region which contains the CLIP sequence.

KW Class II associated invariant chain peptide; CLIP region; MHC;

KW major histocompatibility complex; T cell epitope; antigen processing;

KW multiple sclerosis; insulin-dependent disorder; rheumatoid arthritis;

KW myasthenia gravis; allergy; tumour; virus; parasite; mycobacterium;

infection; transplant rejection; immunostimulant; diagnosis.

Length: 74 February 11, 2000 15:48 Type: P Check: 8373 WILRRLPHVH PPQGASREEP GRLPQGRRRR RQTVLLASIY LASLLPIEIV

20-MAR-1997. 16-SEP-1996; D01763. 14-MAY-1996; US-017663. 14-SEP-1995; DE-034170.

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pr tool - based on primary sequence of invariant chain contg. primary prequence of specific T-cell epitope
ps Disclosure; Page 8; 29pp; German.
CC This is the sequence of amino acids 97-126 from the p35 isoform of the invariant chain (i.e. of a polypeptide which associates with cc major histocompatibility complex class II molecules in the ce endoplasmic reticulum to form a class II molecules in the cc endoplasmic reticulum to form a class II loading compartment).
CC Claimed polypeptides are derived from the invariant chain by cc replacing at least part of the CLIF (CLass II associated Invariant Cc chain peptide) sequence with a T cell epitope. The recombinant cc polypeptide is processed in antigen presenting cells to produce the specific T cell epitope. Such polypeptides are useful in vaccines cc for the treatment of autoimmune disorders, e.g. rheumatoid arthritis, cc multiple sclerosis, insulin-dependent diabetes mellitus and myasthenia cc gravis, allergies and tumours, for the control of viral, parasitic and cc mycobacterial infections, and for use against transplant rejections. CC They can also be used in the diagnosis of autoimmune disorders.
CC Vaccines produced using the recombinant polypeptides have a higher cc immunostimulant specificity and selectivity over prior art vaccines. Sequence 30 AA;
PT DNA encoding prorribosome inactivating proteins - inactive procedures of ribosome inactivating proteins; can be expressed in procedures of ribosome inactivating protein; can be expressed in procedures of ribosome inactivating protein, RIP) protein call 4; Column 117-120; 186pp; English.

CC W25141 shows a Luffin-A (ribosome inhibitory protein, RIP) protein produced which was engineered to contain a selectively removable internal peptide contain was engineered to contain a selectively removable internal peptide contains requant and beta units of the RIP. When CC enkaryotic ribosomes and hence preventing protein production. Many contains the regard activity and are capable of inactivating contains regard and the capacity and the removable internal linker including contains. The RIPs can be used in the construction of therapeutic coxins targeted to specific cells such as tumour cells via the coxins targeted to specific cells such as tumour cells via the coxins targeted to specific cells such as tumour cells via the coxins targeted to specific cells such as tumour cells interest contains targeted to specific cells, pecause of the capacity to provide correct post-translational processing. However, CC antachment of a targeting polypeptide, e.g. a monoclonal antibody. CC RIPs effectively inhibit protein synthesis in eukaryotic cells resulting contains a contain the cells and the contains the contains are not cytotoxic to contain the contains are not cytotoxic to the capacity cells, they can be recombinantly expressed in such cells and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            W13647 Length: 30
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11-JUN-1990; 535636.
09-DEC-1992; US-987927.
11-JUN-1990; US-535636.
11-JUN-1995; US-78761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Luffin-A (a ribosome inhibitory protein) inactive precursor.
Maize; proRIF; ribosome inactivating protein; alpha; beta subunit;
internal linker: Barley Translation Inhibitor; Trichosanthin;
Ricin A-chain; Abrin-A A-chain; Saporin; SLT-1; Luffin A; MAP;
Ricinus communis agglutinin; Momordin; pap-S; Luffin-B; Dianthin 30;
Ricinus communis agglutinin; Momordin; pap-S; Luffin-B; Dianthin 30;
therapeutic toxin; tumour cell trargeted; protein synthesis inhibitor;
post-translational modification; cancer; neoplasia; HIV: AIDS;
human immunodeficiency virus; acquired immune deficiency syndrome.
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W25146

ANVSFSLSGA DSKSYSKFIT

ALRKALPSKE KVSNIPLLLP SASGASRYII

151

TLVTIPYSGN

YERLQNAAGK IREKIPLGFR ALDSALTSIF HYDSTAAAAA

YIMGYLVNST

SYFANESDAK

QLAQTNNGAF

RTPVVIIDNK GQRVEITNLA SKVQIKDVNS KLLLNKQNIP

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II AA_SEQUENCE 1.0

ID W25146; standard; p

AC W25146; criboso

AC W25146; prorr; rib

BE Luffin-B (a riboso

KW Malze; prorr; rib

KW Ricinus communis a

KW Ricinus communis a

KW Ricinus communis a

KW Ricinus communis a

KW Post-translational

KW Numan immunodefici

Synthetic.

PN 08-7UI-1997.

PF 11-7UN-1995; US-98

PR 11-7UN-1997; US-98

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Luffin-B (a ribosome inhibitory protein) inactive precursor.

Maize; proRIF; ribosome inactivating protein; alpha; beta subunit;

Internal linker; Barley Translation Inhibitor; Trichosanthin;

Ricin A-chain; Abrin-A A-chain; Saporin; SIT-1; Luffin-B; Dianthin 30;

Ricinus communis agglutinin; Momordin; PAP-S; Luffin-B; Dianthin 30;

therapeutic toxin; tumour cell targeted; protein synthesis inhibitor;

post-translational modification; cancer; neoplasia; HIV: AIDS;

human immunodeficiency virus; acquired immune deficiency syndrome.
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Length: 250
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US-987927.
US-535636.
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February 11, 2000 15:48
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Type: P Check: 7681
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W22410;

standard;

Protein;

140

æ

08-DEC-1997

Alpha-4 integrin mouse MAb 21.6 VH region.

Alpha-4 integrin; humanised antibody; monoclonal antibody 21.6;

asthma; atherosclerosis; AIDS; dementia; diabetes; tumour;

metastasis; inflammatory bowel disease; rheumatoid arthritis;

transplant rejection; graft versus host disease; nephritis;

atopic dermatitis; psoriasis; myocardial ischaemia;

acute leukocyte mediated lung injury; therapy.

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PT Uses of humanised alpha-4 integrin antibody - for treatment of PT asthma, atherosclerosis, AIDS, dementia, etc.

PT claim 25; Fig 7; 10/pp; English.

CC This polypeptide, designated Ha, comprises the heavy chain variable CC region (VH) of a humanised alpha-4 integrin antibody 21.6. It is CC composed of complementarity determining regions (CDRs) from the VH CC region (see W22410) of mouse alpha-4 integrin monoclonal antibody CC 21.6 and a modified human 21/28 CL framework. It can be expressed in mammalian host cells following PCR amplification and mutagenesis CC of appropriate fragments of mouse and human DNA sequences. The CC humanised 21.6 VH and a humanised 21.6 VL (see W22412) can be used to produce a claimed humanised 21.6 vL (see W22412) can be used CC manufacture of a medicament for treating asthma, atherosclerosis, CC AIDS, dementia, diabetes, inflammatory bowel disease, themustoid CC arthritis, transplant rejection, graft versus host disease, tumour cometastasis, nephrifis, atopic dermatitis, psoriasis, myocardial CC ischaemia, and acute leukocyte mediated lung injury. The antibody comay also be used in the affinity purification of alpha-4 integrin CC for use as a vaccine or an immunogen. It is also useful for generating diotypic antibodies. The humanised antibody has a ccomplement of the human circulation essentially equivalent to that CC of naturally occurring human antibodies.
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29-MAY-1997.
21-NOV-1996; U18807.
21-NOV-1995; US-561521.
(ATHE-) ATENAN NEUROSCIENCES II
Bendig MM, Jones ST, Leger O.
WPI; 97-297879/27.
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W22413;
08-DEC-1997 (first entry)
Humanised alpha-4 integrin antibody; monoclonal antibody 21.6;
Alpha-4 integrin; humanised antibody; monoclonal antibody; as thereosclerosis; AIDS; dementia; diabetes; tumour; metastasis; inflammatory bowel disease; rheumatoid arthritis; transplant rejection; graft versus host disease; nephritis; transplant rejection; graft versus host disease; nephritis; acoute leukocyte mediated lung injury; therapy.
Chimeric Mus musculus; Chimeric Homo sapiens;
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/label= CDR3
/note= "21.6
113. .123
/label= FR4
/note= "21/28
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67. .98
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36. .49
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/note= "21/28'CL
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'note = "21.6
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/note= "21/28'CL framework region
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L, important in supporting the CDR2
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/label= CDR3 /note= "complementarity determining

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iabel= CDR1
note= "complementarity determining

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FR2

region region

/label= 20. .49

Leader

Location/Qualifiers

/label= Fki /note= "framework region

peptide Key

Mus musculus

region

/label=

FR3 CDR2

"framework region

"complementarity determining region

2

WC9718838-A1.
29 MAY-1997.
21 NOV-1996; U18807.
21 NOV-1995; US-561521.
21-NOV-1995; US-561521.
21-NOV-1995; US-561521.
21-NOV-1995; US-561521.
21-NOV-1995; US-561521.
21-NOV-1995; US-561521.
21-NOV-1997879/27.
N-PSDB: 774760.
Uses of humanised alpha-4 inte

OJ,

Saldanha

'n

Yednock

TA;

integrin antibody -

for treatment

of.

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W22410
  Length: 140
 February 11, 2000 15:48
Type: P
Check: 2629
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140 AA;

asthma, atherosclerosis, AIDS, dementia, etc.

Claim 18; Page 69-70; 107pp; English.

Chaim 18; Page 69-70; 107pp; English.

Chaim 18; Page 69-70; 107pp; English.

Complementarity determining regions (CDRs) of the 21.6 The complementarity and a claimed humanised 21.6 The complementary of a medicament for treating a disease selected from asthma, atherosclerosis, AIDS, dementia, diabetes, inflammatory bowel disease, rheumatoid arthritis, transplant rejection, graft versus host disease, tumour metastasis, nephritis, atopic dermatitis, psoriasis, myocardial arthritis, ransplant rejection, graft versus host disease, tumour metastasis, nephritis, atopic dermatitis, psoriasis, myocardial ischaemia, and acute leukocyte mediated lung injury. The antibody may also be used in the affinity purification of alpha-4 integrin for use as a vaccine or an immunogen. It is also useful for invention have a half-life in the human circulation essentially soccurring human antibodies.

!!AA_SEQUENCE 1.0

MAVVTGVNSE VQLQQSGAEL VKPGASVKLS CTASGENIKD

TYIHCVKQRP EQGLEWIGRI DPANGYTKYD PKFQGKATIT ADTSSNTAYL

¹⁰¹ AVYFCAREGY YGNYGVYAMD

W22413

Length: 123

February

11, 2000 15:48

Type: P Check: 9557

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FT W09718838-A1.

PD 29-MAY-1997.

PF 21-NOV-1995; U3-561521.

PR 21-NOV-1995; U3-561521.

PA (ATHE-) ATHENA NEUROSCIENCES INC.

PA (ATHE-) ATHENA NEUROSCIENCE INC.

PA (ATHE-) ATHENAL NEURO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Humanised alpha-4 integrin antibody 21.6 Vt version Ha. Alpha-4 integrin; humanised antibody 21.6 vt version Ha. Alpha-4 integrin; humanised antibody; monoclonal antibody 21.6 asthma; atherosic atros; AIDS; dementia; diabetes; tumour; metastasis; inflammatory bowel disease; rheumatoid arthritis; transplant rejection; graft versus host disease; nephritis; atopic dermatitis; psoriasis; myocardial ischaemia; acute leukocyte mediated lung injury; therapy.
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132. .1
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/label= (
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/label= FR4
/note= "21/28'CL
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"21.6
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"21/28'CL
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W17971 s

standard; 1.0

Protein;

47

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binding protein C-terminal region binding protein; CTBP; modulator; cell proliferation; glycogen.

101

YVSINKLYIS IPSTAANATT

51

NIAANEPEKN MIIDVITSLD

12-DEC 1997 (first entry)
RAC protein kinase C-terminal b
RAC protein kinase C-terminal b
signal transduction; insulin; co
Homo sapiens.
W09718303-Al.
22-MAY-1997.
05-NOV-1995; GB-025704.
15-DEC-1995; GB-023779.

SAON)

W26413

Length: 121

February 11,

2000 15:48

Type: P

Check: 3737

GGCNKCLNIK HAKSQCKPSS

LSPYTLLLDI HFLAQSAFFF THSIVSIALN

LLISSLENVN PGINPRFFIQ

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                                                        against infectious agents, including pseudorables

Example 2; Column 45-48; 70pp; English.

This sequence comprises a polypeptide, designated C22R, encoded this sequence comprises a polypeptide, designated C22R, encoded the sequence strand (open reading frame 3) of the HindIII C by the sense strand (open reading frame 3) of the Shows no fragment (see TM4564) of swinapox virus (SPV). C22R shows no caparent homology to known protein sequences. A claimed recombinant vector comprises a heterologous nucleotide sequence inserted into, or replacing, all or a portion of a non-essential SPV gene or nucleic acid sequence of the HindIII C fragment. The vector can be used for the expression of heterologous proteins, both in vivo as a vaccine, and in vitro for production of the selected protein. The heterologous protein is preferably selected protein. The heterologous protein is preferably to short-restricted to swine, the use of modified recombinant SPV as a live vaccine vector eliminates the risk of spreading infection cath the virus to other animal populations.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    21-APR-1989; 342212.
21-APR-1989; US-908241.
21-APR-1989; US-908630.
21-APR-1994; US-307499.
21-SEP-1994; US-307499.
21-SEP-1996; US-30749.
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a live vaccing the with the virus to 121 /
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Swinepox virus HindIII C encoded protein C
SPV; vaccine; vector; pseudorables virus;
Swinepox virus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   swinepox virus vector - used particularly for vaccines
ectious agents, including pseudorables
    to other animal populations.
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HARVAN SOLUTION OF THE SECOND SOLUTION OF THE
PT DNA encoding chloroplast RecA-like protein from Arabidopsis thaliana PT useful for increasing the frequency of homologous chromosomal recombination when lacking the transit peptide sequence Sclaim 2; Columns 7-9; 14pp; English.

CC The present sequence is of a Rec-A like protein from Arabidopsis CC thaliana. The cDNA encoding this protein (793777) was isolated from an Arabidopsis library screened with a Synechococcus recA gene. The cDNA compared to the transit peptide, is missing. In E. coli, and thus the start of the transit peptide, is missing. In E. coli, and many other prokaryotes, the Rec-A protein is essential for homologous cand many other prokaryotes, the Rec-A protein is essential for homologous that the enzyme encoded by the present sequence may also be concerned with DNA repair. The 5' end of the recA gene, i.e. the portion encoding the transit peptide part of the protein that directs the final product that directs completed proteins to enter the nucleus. By introducing the increase the frequency of homologous recombination in the chromosomal sequence 438 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ---AUG-1992; US-938332.

(CORR) CORNELL RES FOUND INC.

[ Cerutti H, Jagendorf A;

WPI; 97-502391/46.

N-PSDB; T93777.

DNA encodin
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Critis polypeptide comprises the putative C-terminal region of a novel RAC protein kinase C-terminal binding protein (CTBP). Its sequence was deduced from a cDNA clone (T67134) isolated from a clone (T67134) isola
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     _SEQUENCE 1.
W34668 stan
W34668;
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Arabidopsis thaliana Rec-A protein.
Rec-A protein; Synechococcus recA gene; tr
homologous recombination; DNA repair; the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Protein
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Key
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WPI; 97-289279/26.
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/note- "putative chloroplast transit peptide"
51. 438
/label- mature_peptide
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he recA gene.
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W34668 Length: 438 February 11, 2000 15:48 Type: ש Check: 9097

23-JUL-1997. 10-CAN-1997; 100316. 18-JAN-1996; JP-023446. (NAGA-) NAGATA SANGYO CO LTD. (NISS) NISSHIN FLOUR MILLING

G

27-FEB-1998 (first entry)
Novel amylase inhibitor protein 2 subunit isolated from wheat. Amylase inhibitor; wheat flour; wheat; human alpha-amylase; blood sugar level; insulin secretion; fatty acid level; appetis suppressor; treatment; diabetes; prevention; obesity.
Triticum sp.
EP-785214-A2.

appetite

!!AA_SEQUENCE

1.0

standard; peptide; 124

AA.

101 51

LPIVIDASGD GAYVCKGVAA

WCRCGALYSM

SGPWMCYPGY AFKVPALPGC RPVLKLQCNG SQVPEAVLRD CCQQLADISE

LDSMYKEHGV QEGQAGTGAF PSCRREVVKL TAASITAVCK

DSQLVLSLKL NPSFTPLSPL FPFTPCSSFS PSLRFSSCYS RRLYSPVTVY

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C W24491;

T 27-FEB-1998 (first entry)

E Novel amylase inhibitor protein subunit isolated from wheat.

W Amylase inhibitor; wheat flour; wheat; human alpha-amylase;

W blood sugar level; insulin secretion; fatty acid level; appetite

W suppressor; treatment; diabetes; prevention; obesity.

EP-785214-A2.

PN EP-785214-A2.

PP 10-JAN-1997; 100316.

PR 18-JAN-1997; 100316.

PR (NAGA-) NAGATA SANGYO CO LTD.

PA (NAGA-) NAGATA SANGYO CO LTD.

PA (NISS ) NISSHIN FLOUR MILLING CO.

PA (NISS ) NISSHIN FLOUR MILLING CO.

PI Miyazaki T, Morimoto T, Murayama R;

PT New isolated proteins from wheat active as amylase inhibitors - used

PT in the treatment and prevention of diabetes and in the prevention of
      W24491
                              claim 1; Page 16; 22pp; English.

This sequence represents a novel amylase inhibitory protein which is composed of two subunits, each having the present sequence. This protein can be prepared by extraction from wheat flour, wheat or wheat gluten. The protein is spherical and has S-S bonds, and has a high inhibitory activity against human alpha-amylase. It has about 5-30 times higher inhibit increases in blood sugar levels, inhibit insulin secretion and inhibit an increase in fatty acid levels, inhibit insulin secretion and feeling of satiety after meals, and easily suppressing appetite. The provention of obesity.

Sequence 122 AA;
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    Length:
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    122
February 11,
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2000 15:48
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BAB
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                                                                                                                                                                                                            W24493 Length: 124
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This sequence represents a novel amylase inhibitory protein which is composed of two subunits, each having the present sequence. This protein can be prepared by extraction from wheat flour, wheat or wheat gluten. The protein is spherical and has s.s. bonds, and has a high inhibitory activity against human alpha-amylase. It has a higher inhibitory activity as compared with several known amylase inhibitors. It can also inhibit increases in blood sugar levels, inhibit insulin secretion and feeling of satiety after meals, and easily suppressing appetite. The proteins can be used in the treatment and prevention of diabetes and the provention of obesity.
                                                                                                                                                                                                                                         Claim 5; page 18; 22pp; English.

This sequence represents a novel amylase inhibitory protein which is composed of two subunits, each having the present sequence. This protein can be prepared by extraction from wheat flour, wheat or wheat gluten. The protein is spherical and has 5-5 bonds, and has a high inhibitory activity against human alpha-amylase. It has about a higher inhibitory inhibit increases in blood sugar levels, inhibit inscreases in can also inhibit an increase in fatty acid levels in the blood, thus maintaining if celling of satiety after meals, and easily suppressing appetite. The proteins can be used in the treatment and prevention of diabetes and the prevention of obesity.

Sequence 124 AA;
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18-JAN-1996; JF-023446.

(NAGATA SANGYO CO LTD.

(NISS ) NISSHIN FLOUR MILLING CO.

MIYSZAKI T, MOLIMOTO T, Murayama R:

MPI; 97-365900/J4.

New isolated proteins from wheat active as amylase inhibitors - used in the treatment and prevention of diabetes and in the prevention of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-FBB-1998 (first entry)
Novel amylase inhibitor protein 3 subunit isolated from wheat. Novel amylase inhibitor; wheat flour; wheat; human alpha-amylase; blood sugar level; insulin secretion; fatty acid level; appetite suppressor; treatment; diabetes; prevention; obesity.
Triticum sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Miyazaki T. Morimoto T. Murayama R: \text{WPI}; 97-365900/34. WPI; 97-365900/34 from wheat active as amylase New isolated proteins from wheat active as amylase in the treatment and prevention of diabetes and in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE 1.0
W24493 standard; peptide; 124 AA
w29523 standard; Protein; 124
w29523;
09-MAR-1998 (first entry)
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                                                                                                                                    WCRCGALYSM
                                                                                                                                                                       SGPWMCYPGQ AFQVPALPAC
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the prevention
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The protein comprises the subunit of the wheat amylase inhibitor from wheat - for inhibiting the claim 2; Page 16; 22pp; English.

Claim 2; Page 16; 22pp; English.

Claim 2; Page 16; 22pp; English.

Chis protein comprises the subunit of the wheat amylase inhibition of or comprises the subunit of the wheat amylase inhibiting an accumulation of visceral fat and hence for preventing visceral fat obesity is claimed. It contains an amylase inhibitor of wheat corigin as an active ingredient. The amylase inhibitor is selected from 0.26 Ala, 0.25 Alb (see W29524) and 0.19 Al (see W29525). The 0.26 Ala protein has inhibitory activity of 26,100 U/mg against human pancreatic alpha-amylase. Also claimed is a food additive comprising 0.26 Alb. The amylase inhibitors can inhibit the capture in sulin secretion and reduce the activity of lipogenic enzymes in the viscus.

Sequence 124 AA;
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                                            11-JAN-1997; 100399.

11-JAN-1997; JP-023445.

A (NAGA-) NACATA SANCYZ CO LITD.

A (NISS ) NISSHIN FLOUR MILLING CO.

BY (NISS ) NISSHIN FLOUR MILLING CO.

A (NISS) NISSHIN FLOUR MILLING CO.

A (NIS) NISSHIN FLOUR MILLING CO.

A (NIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W29524 S
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Amylase inhibitor; 0.26 Ala; wheat; visceral fat; obesity; therapy.
Triticum aestivum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (NAGA-) NAGATA SANGYO CO LTD.
(NISS) NISSHIN FLOUR MILLING CO.
Godd T, Miyazaki T, Morimoto T,
WPI: 97-365735/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              09-MAR-1998 (first entry)
Wheat amylase inhibitor 0.26 Alb subunit.
Amylase inhibitor; 0.26 Alb; wheat; visceral
enzymes in
Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SGPWMCYPGY AFKVPALPGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; Protein; 122
   the vis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP-023445.
                         viscus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LDSMYKEHGV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               February 11, 2000 15:48 Type: P Check: 2685
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murayama R,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    fat; obesity; therapy
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W29524

Length: 122

February

11,

2000 15:48

Type:

SGPWMCYPGY WCRCGALYSM

AFKVPALPGC

RPVLKLQCNG SQVPEAVLRD

CCQQLADISE

LDSMYKEHGV

QEGQAGTGAF

PSCRREVVKL TAASITAVCK

LPIVIDASGD GAYVCKGVAA

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PF 11-JAN-1996; JP-023445.

PR 18-JAN-1996; JP-023445.

PR (NISS) NISSHIN FLOUR MILLING CO.

PA (NISS) NISSHIN FLOUR MILLING CO.

PA (NISS) NISSHIN FLOUR MILLING CO.

PA (NISS) NISSHIN FLOUR MILLING CO.

PI GOGA T, Miyazaki T, Morimoto T, Murayama R, Takase S;

PH WPI: 97-365735/34. Inhibitor from wheat - for inhibiting the

PT accumulation of visceral fat to prevent visceral fat obesity

PS Claim 2; Page 18; 22pp; English.

C Inhibitor homodiner protein 0.19 Al. An agent for inhibiting an

CC accumulation of visceral fat and hence for preventing visceral

CC Inhibitor homodiner protein 0.19 Al. An amylase inhibitor of wheat

CC origin as an active ingredient. The amylase inhibitor is selected

CC origin as an active ingredient. The amylase inhibitor of vheat

CC oungrishing 0.26 Alb and 18 (see W39254) and 0.19 Al. The

CC comprishing 0.26 Alb. The amylase inhibitors can inhibit the

CC dispertion of starch, inhibit increases in blood glucose level,

CC enzymes in the viscus.

Sequence 124 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 !!AA_SEQUENCE 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               W29525
         Polynucleotide(s) encoding type I ribosome-inactivating proteins which are suitable for use as components of cytotoxic therapeutic agents.

Disclosure; Fig 4; 221pp; English.

R63906 is the type I ribosome-inactivating protein (RIP) 14 fin one of the nine RIPs described in R63903-R63911.

RIPS are key components of cytotoxic therapeutic agents (CTAs), which include gene fusion products and immunoconjugates. CTAs may be used to selectively eliminate any cell type to which a RIP component is targetted, by the specific binding capacity of the second component of the agent. They can used in the treatment of diseases where the elimination of a particular cell type is desired, such as autoimmune disease, cancer and graft-versus-host
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Type I ribosome inactivating protein luffin, RIPs;
Type I ribosome inactivating proteins; luffin, RIPs;
cytotoxic therapeutic agents; autoimmune disease; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      101 LPIVVDASGD GAYVCKDVAA YPDA
                                                                                                                                                                                                                                                                          12-MAY-1993; US-064691.
(XOMA ) XOMA CORP.
Better MD, Carroll SS,
WPI; 95-006804/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-MAR-1998 (first entry)
Wheat amylase inhibitor 0.19 Al subunit.
Amylase inhibitor; 0.19 Al; wheat; visceral fat; obesity; therapy
Triticum aestivum.
                                                                                                                                                                                                                                                                                                                                                                                                                                        graft-versus-host disease
                                                                                                                                                                                                                                                                                                                                                                                                 w09426910-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-JUL-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              51
                                                                                                                                                                                                                                                                                                                                                              2-MAY-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Protein; 248 AA
                                                                                                                                                                                                                                                                                                  Studnicka GM, Carroll SF;
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IIAA_SEQUENCE 1.0

ID W45093; standard; peptide; 176 AA.

AC W45093;

DT 07-MAY-1998 (first entry)

DT 07-MAY-1998 (first entry)

DT 07-MAY-1998 (first entry)

DT 07-MAY-1998 (first entry)

Residues 217-392 of human type B EBNA2 (strain AG876).

Residues 217-392 of human type B EBNA2; transplant.

SEPT Region 69.76

PR 10-1997 (Jabel Conserved_region_6

PN W09730081-A1.

PD 21-AUG-1997 (Jabel Conserved_region_6

PN W09730081-A1.

PD 21-AUG-1997 (Jabel Conserved_region_6

PN W1-197-424974/39.

PR W1-197-42497/39.

PR W1-197-42497/
I!AA_SEQUENCE 1.0

ID W38190 standard; Protein; 107 AA
AC W38190;
DT 11-MAY-1998 (first entry)
DE Soybean SCARECROW SRP91 protein.
KW SCARECROW, SCR gene; SRP91; tran
KW crop improvement; soybean.
OS Glycine max. Location/Qualifi
FH Key Location/Qualifi
FT Misc_difference 23
FT W09741152-A1.
PD 06-NOV-1997;
DF 25-APR-1997; U707022.
PR 26-APR-1997; US-842445.
PR 26-APR-1995; US-84845.
PR 26-APR-1995; US-848617.
PA (UYNY) UNIV NEW YORK STATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              useful to provide protection against Epstein Barr virus Disclosure; Fig 5; 62pp; English.

The patent discloses five new peptides which display substantial homology with the CBFI/RBDJK interaction domain of EBNA2 (see M24280-W24284). In particular, the peptides include amino acids 318-327 of EBNA2 (i.e. conserved region 6, or CR6). The peptides, when administered in an amount sufficient to inhibit interaction of EBV EBNA2 with CBFI, are used to protect mammals prior to or following a transplant, against Epstein Barr virus. Labelled peptides are used to detect factors that interact with EBV EBNA2. The present sequence is from one strain of EB virus (human type B used to identify CR6 (GPPWMPPV) residues 69-76).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                W09730081-A1.
21-AUG-1997; U02243.
12-FEB-1996; US-011557.
13-FEB-1996; US-011557.
(UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE.
(UYJO ) UNIV JOHNS HOPKINS SCHOOL MEDICINE.
Peptide with homology to CBF1-RBPJk interaction domain of EBNA2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                            11-MAY-1998 (first entry)
SOybean SCARECROW SRP91 protein.
SCARECROW, SCR gene; SRP91; tran
crop improvement; soybean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HQLSLPPHPP PHQSTPHCSS DSTGLPPPPT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TLSNYDGKAI TVAVDVSQLY IMGYLVNSTS YFFNESDAKL
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                                                                                                                                                                                                               Location/Qualifiers 23 /note- "encoded by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GRGRMHKLPE PRRPGPDTSS
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                                                                                                                                                                                                                   "encoded by NNC
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                                                                                                                                                                                                                                                                                                                                                                                       transgenic plant;
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type B)
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n В,

an be

antibiotic used as used as animal feed

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PT Streptomyces frenolicin yells (specially actions) actili protein. CC This is a ketoreductase related to the actinorhodin actili protein. CC It is the encoded product of gene P of a claimed Streptomyces claimed coding sequences (genes A-U) that respectively encode 21 CC proteins (see W34199-219) involved in frenolicin synthesis. The 21 CC genes of the cluster can be divided into 5 subclusters: (1) genes CC A, B, C, D and F encode an efflux pump; (2) genes H, I, J and K CC polyketide synthases; (4) genes (9) of and R encode a hemiketalase, CC polyketide synthases; (4) genes (9) of and R encode a hemiketalase, CC encode a keto/encyl reductase and hydrolase. Also claimed are created a keto/encyl reductase and a hydrolase. Also claimed are Streptomyces roseofulvus). Host cells can be cultured to produce the colls (especially a Streptomyces can be cultured to produce the colls (especially a Streptomyces can be cultured to produce the colls (especially a Streptomyces can be cultured to produce the colls (especially a Streptomyces can be cultured to produce the colls (especially a Streptomyces can be cultured to produce the colls (especially especially 
                                               W38190
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02-MAY-1997; 107329.
07-MAY-1996; US-016753.
(HOFF) HOFFMANN LA ROCI
Reeves CD, Soliday CL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       controls gravitropic response in aerial structures and roct formation. SCR proteins (see W38178-201) from dicots and monocots, SCR proteins lacking 1-4 of MOTIFF I-VI, and polypeptides corresponding to MOTIF I, II, III, IV, vor VI of SCR are claimed. Transgenic plants can be insineered to overexpress the SCR protein, so that cell division is increased in roots, resulting in thicker root development, while a plant with an altered stem or hypocotyl gravitropism is less susceptible to lodging. Plants that contain an antisense molecule that suppresses the expression of endogenous SCR gene product have thinner root development. A gene of interest plant to confer herbicide, salt, pathogen or insect resistance, or the second of the second of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 97-538619
N-PSDB; T93095
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               11 MAY-1998 (first entry)
Streptomyces ketoreductase (frenolicin gene P product).
Frenolicin; antibiotic; feed additive; anticoccidial;
coccidiostatic; ketoreductase; Streptomyces roseofulvus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SRPg1 (see T95765). It shows homology to Arabidopsis SCARECROW (SCR) protein (see W88178). SCR is a member of a novel protein family and contains a number of potential functional domains similar to those found in transcription factors. SCR is expressed specifically in embryo root progenitor tissue and in certain root and stem tissues. It regulates as specific asymmetric division, and controls gravitropic response in marrial asymmetric division, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               101
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This is the deduced amino acid sequence of soybean genomic SRPg1 (see 195765). It shows homology to Arabidopsis SCARE (SCR) protein (see W38178). SCR is a member of a novel pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Streptomyces sp.
EP-806480-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE 1.0
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Sequence 107 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   51
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WPI; 97-549683/50.
N-PSDB; T95765.
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Pysh L, Wysocka-Diller
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root development,
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alter plant
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18-DEC-1990; 79-412020.

(GREC) GREEN CROSS CORP.

(GHIM/) SHIMOTONO K.

WPI; 93-260858/33

Protein contg. non-A non-B hepatitis antigen fragment - prepd. by culturing transformants transformed by vector contg. base sequence coding specified aminoacid sequences, used for detecting hepatitis claim 1; Fig 13; 53pp; Japanese.

The present sequence is a non-A non-B hepatitis virus (NANBH) or hepatitis C virus (HCV) antigen, useful for diagnosis or detection. Sequence 239 AA;
                                                                                                           14 - J;
23 - D]
                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  101
                                                                                                                                                                                                                                                                                                                  11-JUN-1998 (first entry)
HRSV protein 22K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       201
                                         (PHAA)
                                                                                                                                                                                                                               uman respiratory syncytial 
JS5716823-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     non A non-B hepatitis virus; NANBH; hepatitis C virus; antigen; diagnosis; detection.
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Hepatitis C virus antigen.
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                                                                                                                                                                                                                                                                                              glycoprotein F;
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             PHARMACIA & U
PL, Wertz GW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard;
                                                            988; US-218737.
986; US-818740.
986; WO-U02756.
992; US-897171.
997; US-854783.
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                                                                                                                                                                                                                                                                                         gpF; glycoprotein G; gpG;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (NANBH) or or detection.
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98-144802/13

N-PSDB; V18737.

Production of human respiratory syncytial virus glyco-protein F or G
- by culturing eukaryotic host cells transfected with corresponding DNA

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REGERENCE COCCONTRACTOR
                                                                                                                                                                   W50006
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15-JAN 1998
08-JUL-1997; U11815.
09-JUL-1996; US-677206.
(HARD ) HARVARD COLLEGE.
Dowhanick JJ, Howley PM;
WPI; 98-100815/09.
N-PSDB; V18818
                                                                                                                                                                                                                                                                                                                                                                 Treating papilloma virus infections, by repressing E6 and E7 oncogenic proteins - using E2 polypeptide or nucleic acid encoding it, useful for treating warts and cervical cancer claim 22; Pages 51-53; 71pp; English.

The present sequence is the human papillomavirus-18 (HPV-18) E2 protein.
                                                                                                                                                                                                                              protein, comprising a DNA-binding and transcription activating domain, or the gene encoding the E2 protein to infected cells. The method is applicable to ilvestock, zoo animals, pets and humans, specifically for the treatment, prevention or reversal of warts, e.g. plantar, common, Butcher's common, flat or genital warts, or epidermodysplasia verruciformis, or growth of papillomavirus transformed cells, especially laryngeal papilloma or cervical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         151
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The present sequence was used in the development of a novel method for the production of human respiratory syncytial virus (HRSV) glycoprotein F (gpF) or glycoprotein G (gpG). The method comprises culturing eukaryotic host cells transfected with an isolated DNA sequence encoding HRSV gpF or gpG. The gp can be used to prepare vaccines against HRSV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         II-JUN-1998 (first entry)
Human papillomavirus-18 E2 protein.
Human papillomavirus-18; HPV-18; E2 protein; infection; tre
wart; epidermodysplasia verruciformis; laryngeal papilloma;
cervical carcinoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Key
                                                                                                                                                                                                                                                                                                                                                     Papillomavirus infections can be treated by administering
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human
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                                                                                                                                                              Length: 365 February 11, 2000 15:48
NDSMCSTSDD TVSATQLVKQ LQHTPSPYSS TVSVGTAKTY GQTSAATRPG
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                                 DKTATCVSHR GLYYVKEGYN
                                                                                                                                MOTPRETLSE
                                                                TCEELWNTEP
                                                                                                 REHGIQTLNH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       papillomavirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        standard;
                                                                                                                                                                                                   365 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "-
                                                                                                                                RLSCVQDKII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
1. .203
/note= "transcription activating domain"
282. .365
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                                                                                               QVVPAYNISK
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                               TFYIEFKSEC
                                                                                                                                DHYENDSKDI DSQIQYWQLI
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                                                                                               MALQGLAQSR
                               EKYGNTGTWE VHFGNNVIDC
                                                                NCMTYVAWDS
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                                                                                                                                                                  Type: P
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PT Helicobacter pylori nucleic acid sequences and encoded PT Helicobacter pylori nyesful in vaccines to treat or prevent H. pylori infection and for diagnosis of H. pylori infection PT infection and for diagnosis of H. pylori infection PT infection and for diagnosis of H. pylori infection PT infection and for diagnosis of H. pylori infection or to identify PT PT infection or to identify H. pylori CC Infis sequence is a H. pylori cell envelope outer membrane protein (OMP) CC having a C-terminal tyrosine cluster motif The protein may be used in a cycline to prevent or treat H. pylori infection or to identify H. pylori pylopeptide binding compounds, useful as potential H. pylori life Cycle activators or inhibitors. The DNA and probes derived from it may be compounded to the pylori in a sample and the diagnosis confidence of H. pylori infection. Nucleic acid sequences complementary to the DNA ct as antisense sequences and can be used to prevent the translation of the pylori mRNA. Antibodies against the protein can be used in immunoassays to evaluate the abundance and distribution of CC (ATCC 55679) was determined from overlapping contils generated by CC mechanically shearing the bacterial DNA. The sequences were analysed for ORF of at least 180 nucleotides, and the predicted coding regions confidence development, the amino acid sequences predicted coding regions for vaccine development, the amino acid sequences predicted from various ORF were analysed for significant homology to other known or exported membrane proteins. Having identified and determined the sequences of interest, particular regions can be isolated from H. pylori by PCR amplification for recombinant polypeptide production, e.g. in E. coli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (ASTR ) ASTRA AB.
Alm RA, Smith D;
WPI; 97-503122/46.
N-PSDB; V24999.
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02-APR-1996;
25-OCT-1996;
28-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cytoplasmic; vaccine; previdentification; binding coinhibitor; duodenal ulcer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE 1.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLRYRLRKHS DHYRDISSTW HWTGAGNEKT GILTVTYHSE TQRTKFLNTV
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Protein; 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24.429510_c2_46.aa cell envelope OMP.
prevention; treatment; infection; envelope;
ng compound; bacteria; life cycle; activator;
lcer disease; chronic gastritis; diagnosis.
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W55590
 Length:
February 11, 2000 15:48
Type: P
 Check: 428
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Sequence

307 AA;

- 301 51 IDRYGDENYL DLLINWINDK NELSLGYKYF ALNQAINNAK NSLFPNTKAI RDVQNALNAV MKKVLLLTLS LSLSEWLHAE RNGFYLGLNF LGASTWFASN RASQEYVERR LGKKRIIGFR GGSSVELGVK VPAFKVNYYS DDYGDKLDYK RVVSVYLNYI HSLFFGYQLG VKGLSIFYKD KDSNKIANRF AGNGGSGGIF AEGSYIQGQG SIGEKASAEN RTIFQLQGKF GVRFSSDEYD GVGSVPGSGL MTGRTLDADT LKRASRHIIR IAFLPYGENT
- !!AA_SEQUENCE 1.0
 ID W55815 standard; Protein; 271

infection

and therapeutic

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IIAA_SEQUENCE 1.0

ID W56137 stand
AC W56137,
AC CALLERY AC ALLERY AC ALLE
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11094395-A.

11-APR-1998.

07-MAY-1997; 116652.

04-APR-1997; US-042935.

07-MAY-1996; US-016753.

(HOFF) HOFFMANN LA ROCHE & CO AVERT, 98-279231/25.
                                                                                                                                                                                                                                                                                                                                                                                                              21-APR-1998.
30-NOV-1995; 565386.
30-NOV-1995; US-565386.
(UYRP ) UNIV ROCHESTER.
Bavoil PM, Hsla R;
WPI; 98-260507/23.
N-PSDB; V16865.
Bacteriophage phi-CPG1 genomic DNA - is useful for transforming mammalian Chlamydia strains
Claim 12: Columns 23-24: 28pp; English.
The present sequence represents the product of the second open reading frame of a circular bacteriophage, designated phi-CPG1. This bacteriophage is able transfect Chlamydia psittact strain Guinea pig Inclusion Conjunctivitis (GPIC). Phi-CPG1 was identified by homology to a previously characterised Chlamydia bacteriophage of avian C. psittaci named Chp1. Chlamydia strains can be responsible for blindness through occular chronic disease trachoma. Phi-CPG1 can be used as a cloning vector, especially as a means to transfer mutant or foreign genes into Chlamydia. Phi-CPG1 can also be used to develop Chlamydia vaccines,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14-JUL-1998 (first entry)
Open reading frame 2 peptide of bacteriophage phi-CPG1.
Circular; bacteriophage phi-CPG1; avian C. psittaci; Chp1;
Chlamydia psittaci strain Guinea pig Inclusion Conjunctivitis;
blindness; ocular chronic disease trachoma; vaccine.
Bacteriophage phi-CPG1.
BS5741697-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prenolicin gene cluster - useful as an antibiotic Claim 8; Page 30; 50pp; Japanese.

The present sequence represents a protein from the frenolicin gene cluster from Streptomyces roseofulvus. The present invention describes: (1) a vector containing the frenolicin gene cluster DNA sequence operably connected to an expression control sequence; (2) a host cell transformed by the above vector; (3) a protein coded by the above DNA sequence; (4) a method for the preparation of frenolicin or a present of the above cell is cultured and frenolicin or its biosynthetic intermediate for it in which the above cell is cultured and frenolicin or its biosynthetic intermediate is isolated from the culture or the cell; (5) a method for the preparation of frenolicin B by oxidising frenolicin, and (6) a method for the preparation of a frenolicin by mixing frenolicin with other components.
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15-JUL-1998
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271 AA;
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roseofulvus frenolicin gene cluster protein 16.
roseofulvus; frenolicin gene cluster; frenolici
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OKKWEDAC

W28052 standa

standard;

Protein;

269

B

251

PVYLALLPPD VTVLSRIHAR

AEGPHGQFVS

EKRVEQW

KLSEQRKGDK ILLNACCPGW

VRTDMAGPKA TKSPEEGAET

101

TPFHIQAEVT QAEGLSPRFH

MKTNFFGTRD VCTELLPLIK PQGRVVNVSS

51

MSSGIHVALV TGGNKGIGLA IVRDLCRLFS

QLDIDDLQSI

RALRDFLRKE YGGLDVLVNN

AGIAFKVADE TRGQAAVQQL

GDVVLTARDV

151

ETITEEELVG

LMNKFVEDTK KGVHQKEGWP

SSAYGVTKIG

w28652;
27-ANG-1998 (first entry)
Anino acid sequence of phospho-beta-glucosidase (bglA).
Anino acid sequence of phospho-beta-glucosidase (bglA).
Staphylococcus aureus protein; ribozyme; antisense sequence; contro.
Staphylococcal gene; regulatory element; bacterial gene expression; vaccine; Staphylococcal infection; food poisoning; scaled skin synd toxic shock syndrome.

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ន្ត្រព្ឋ
                          PA (INCY) INCYTE PHARM INC.

PA (INCY) INCYTE PHARM INC.

PR GOAL SK, Hillman JL;

DR WPI, 98-321527/28.

PT DNA encoding human carbonyl reductase polypeptide - useful for recombinant production of the enzyme and anti:sense treatment of PT diseases associated with the enzyme and increase protein CC (INCA) sequence. Also claimed are: a probe that hybridises to the DNA (INCR) sequence. Also claimed are: a probe that hybridises to the DNA (INCR) sequence; an expression vector containing the bukh sequence; and a host CC sequence; an expression vector containing the production of CC carbonyls to alcohols, physiologically functioning as dehydrogenases of CC crasting and nor hydroxysteroids, as well as in drug metabolism. The cell CC can be used to produce a recombinant human carbonyl reductase protein, CC which can be used to treat inflammatory or immunological disorders CC development after exposure to carcingens. The vector can be used for the CC same purpose. The probe can be used to detect carbonyl reductase nucleic crasting from purpose and be used to detect carbonyl reductase nucleic present sequence represents human liver carbonyl reductase nucleic crasting from the used to control expression of the gene. The CC sequence represents human liver carbonyl reductase.
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    W51011
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26-MAY-1998.
09-DEC-1996;
09-DEC-1996;
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reagents an
Sequence
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Human carbonyl reductase; HCRD; inflammatory disorder; cancer;
immunological disorder; prostaglandin E.
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Length: 277
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ic tests and the study for Chlamydia and strategies in infections. 186 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                      762129.
US-762129.
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    Check: 7216
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W28052
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CC The present sequence represents a Staphylococcus aureus protein, that, based on homology with a Bacillus subtilis protein, is believed to be a seed on homology with a Bacillus subtilis protein, is believed to be a phospho-beta-glucosidase (bglA). The DNA sequence was isolated from a library of clones of S. aureus WCH 29 in Escherichia coli. The DNA sequence can be used in the construction of ribozymes and antisense sequences to control the expression of Staphylococcal genes. The DNA sequences to control the expression of Staphylococcal genes. The DNA control of bacterial gene expression. The produce protein may be used to produce vaccines to enable a host to produce specific antibodies with antibacterial action. These vaccines and antibodies would protect a host against invasion by S. aureus, and conditions relating to skin syndrome, and toxic shock syndrome.
                                                                              02-ARR-1998: 017154.
24-SEP-1996; US-028363.
11-OCT-1996; US-0286603.
24-SEP-1996; US-026603.
(LARB-) LAR BIOTECHNOLOGY INC.
Melkonyan H, Umansky S;
WPI: 98-330704/20.
N-PSDB; V19114.
 New secreted apoptosis-related proteins - useful for mapoptosis, particularly for treatment of prostatic or also for diagnosis and monitoring of disease claim 1; Page 56-58; 101pp; English.

This polypeptide comprises human secreted apoptosis-re
                                                                                                                                                                                                                                                                                                                Human secreted apoptosis related protein hSARP3. Secreted apoptosis related protein; SARP; hSARP3; human; prostate cancer; breast cancer; diagnosis; gene therapy.
                                                                                                                                                                                                                                                                                                                                                                                                   _SEQUENCE 1.0
W37816 standa
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                                                                                                                                                                                                                                            Protein
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(SMIK ) SMITHKLINE BEECHAM CORP.
Black MT, Burnham MK, Hodgson JE, Knowles DJC,
Pratt JM, Reichard RW, Rosenberg M, Ward JM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel polypeptide(s) from Staphyloc to isolate antimicrobial compounds,
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Misc_difference
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                                                                                                                                                                                                                                                                                                                                                                                              standard; Protein; 317
                                                                                                                                                                                                                      Location/Qualifiers
1. 28
/label= Sig_peptide
29. 317
/label= Mat_protein
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1. .269
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "residues designated X are the specification"
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secreted apoptosis-related
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!!AA_SEQUENCE 1.0
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Bacterial growth inhibiting peptide(s) enterocin 900 or brochocin used to inhibit growth of susceptible bacteria in e.g. foodstuff ilving animal, food preparation area or fermentation vessel Example 7; Page 138-139; 174pp; English.

This polypeptide comprises the enterocin-900 pre-protein of Enterococcus faecium 900. Enterocin-900 is a novel broad spectrum bacteriocin that has activity against other strains of Enterococcus as well as many other organisms. The invention includes a method for inhibiting pathogenic bacteria by providing a bacteriocin selected from enterocin-900 and brochocin-C (see W59021), either as a composition or by providing a bacterial source of the bacteriocins. This is used to inhibit spoilage
                                                                                                                                                                                                                  W09809639-A1.

12-MAR-1998.
05-SEP-1997; U15609.
05-SEP-1996; US-026257.
(UYAL-) UNIV ALBERTA.
FIRAT CM, Greer GG, Leisner JJ, McCormick JK, McMullen LM, PRDS J, Poona, Stiles ME, Van Belkum MJ, Vederas JC, Worobo RW, WOT, 98-193319/17.
WPSDB; V1704-05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Enterocin 900; bacteriocin; signal peptide; protein secretion; preservative; food spoilage; lactic acid bacterium; antibacterial. Enterococcus faecium 900.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
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19. .71
/label= Mat_protein
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productining genes for glycerol-3-phosphate dehydrogenase and-or
property glycerol-3-phosphate genes for glycerol-3-phosphate
ps Claim 9; page 36-37; 57pp; English.

Chaim 9; page 36-37; 57pp; English.

Conversion of dihydroxyacetone phosphate to glycerol-3-phosphate.

Chaim 9; page 36-37; 57pp; English that catalyses the

Corganisms that express G3PDH and/or glycerol-3-phosphatase (G3P)

Chaim 9; page 36-37; 57pp; English

Chaim 9; page 36-37; page 
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W57330 stand
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                                      18-SEP-1998 (first entry)
Flea serine protease inhibitor variable domain protein pfSPI14-137
Serine protease inhibitor; SPI; anti-haematophagous ectoparasite;
vaccine; control; flea; acarina; immune response.
Ctenocephalides felis.
W09820034-A2.
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Glycerol-3-phosphate dehydrogenase gpsA.
Glycerol-3-phosphate dehydrogenase; G3PDH; gpsA;
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14-MAY-1998.
05-NOV-1997; U20678
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10-NOV-1997;
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1996; US-030602.
DU PONT DE NEMOURS & CO
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IIAA_SEQUENCE 1.0
IID W55052 standar
AC W55052
DT 28-SEP-1998 (
DE Sunflower anti
KW Antifungal; fix
Antifungal; fix
Antifungal; fix
W Glucose oxidas
KW Grop protectif
OS Helianthus ann
EH Key Cifferen
FT Misc_differen
FT Misc_differen
PN W09813478-A2.
PD 02-APR-1998;
PR (4-SEP-1996; F
PR (4-SEP-1996; F
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Claim 5; Page 53-54; 139pp; English.

This polypeptide is the deduced product of a DNA sequence (see v2725) amplified from sunflower cv. zebulon genomic DNA using PCR primers (see v27249-50) based on tryptic peptides (see w55050-51) of the isolated sunflower w53 antifungal protein.

R Southern blot of sunflower genomic DNA using this DNA fragment revealed the existence of multiple homologous sequences in the sunflower genome. A full-length clone (see v27260) for sunflower sunflower genome. A full-length clone (see v27260) for sunflower m559 (see w55053) was subsequently obtained. Claimed antifungal proteins, which may include this M559 polypeptide sequence, have a mol.wt. of 55-65 kDa (SDS-PAGE), have carbohydrate oxidase (especially glucose oxidase) activity, show anti-Phytophthora and/or anti-Pythium activity, can be expressed in transgenic plants to reduce susceptibility to infection by fungi, or expressed in host cells for use in antifungal compositions. Plants treatments with fungicides and have a longer shelf life.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       04-SEP-1997; E04923.
19-MAR-1997; EP-200831.
04-SEP-1996; EP-202466.
(MOGE-) MOGEN INT NY.
Custers JHHV, Lageweg W.
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The present sequence, designated pfSp114-137, represents Ctenocephalides felis serine protease inhibitor (Sp1) variable domain protein. The Sp1 protein represents a novel target for anti-haematophagous ectoparailtic vaccines and drugs. Compositions containing the inhibitor are used to control fleas (adults or larvae) on mammals or birds, particularly cats and dogs. More generally the composition is also used for control of insects and acarina, on animals or on stored goods, plants, trees etc. including domestic pests and insects that are vectors of disease. The Sp1 proteins are able to induce a protective immune response against Sp1. Sequence 137 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jucose oxidase; transgenic plant; Phytophthora; Pythium; crop protection; disease resistance. Helianthus annus cv. zebulon
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(HESK.) HESKA CORP.
(HESK.) MAGDUX JD, Silver GM, Wisnewski
WPI; 98-286864/25.
N-PSDB; V38773.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           usuers JHHV, Lageweg W, Melchers LS, Ponstein AS, Sela-Buurlage MB, Stuiver MH, Van Deventer-Troost WPI; 98-230692/20.
N-PSDB: V77767
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28-SEP-1998 (first entry)
Sunflower antifungal protein MS59
Antifungal; fungicide; MS59; sunfl
glucose oxidase; transgenic plant;
glucose oxidase; transgenic plant;
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Claim 5; Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New plant proteins having anti-fungal activity - useful as, carbohydrate oxidase(s) for protection against Phytophthora
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W55052

Length:

February

11,

2000

15:48

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4306

W54728 standard; W54728; 25-SEP-1998 (fin

peptide;

24

8

Mannose; antigen;

antigen-presenting cell; mannosylated peptide; T cell;

treatment.

26-SEP-1997; NL0536.
26-SEP-1996; EP-202701.
(UYLE-) RIJKSUNIV LEIDEN.
Drijfhout JW, Koning F;
WPI; 98-230631/20.
Increas*

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PNA encoding v28 seven transmembrane receptor polypeptide - useful for producing recombinant polypeptide and anti-V28 antibodies, and in screening assays for v28 agonists and antagonists.

Claim 1: Columns 61-64: 56pp; English.

The present sequence claimed by the invention represents the human v28 years transmembrane (7TM) receptor. It is encoded by the full length v28 genomic DNA (V18343) or by the v28 CNA (V18352). As the v28 protein (W48722) is a 7TM receptor it is probably involved in signal transduction. The invention also claims that cells transformed with v28 DNA can be used to produce the recombinant polypeptide, to produce anti-v28 antibodies or in screening assays for v28 agonists or antagonists. The antibodies, agonists and antagonists could then be used to modulate v28 receptor-ligand binding, for e.g. in immunological and/or inflammatory events in vivo.
                                                                                                                                                                     W48722
                                                                                                                                                                                                                                                                                                                                                                          17-NOV-1993; 153848.
17-NOV-1992; US-977452.
(ICOS-) ICOS CORP.
GOdÍSKA R, GTAY PW. SCHWE
WPI: 98-32113/29.
N-PSDB: V18343, V18352.
                                                                                                                                                                                                                                                                                                                                                                                                                                     US5759804-A.
02-JUN-1998.
17-NOV-1993;
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W48722 standa
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V28; place
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hmmunology; inflammation.
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                                                                                                MCKETTAFFF
                                                                                                                       LVVFALTNSK
                                                                                                                                               MDQFPESVTE NFEYDDLAEA CYIGDIVVFG
                         IFLETLKLYD
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                         LIYAFAGEKF
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                                                                                                RTVQHGVTIS
                                                                                                                                               IFAIGLVGNL
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!!AA_SEQUENCE 1.0

ALLLL

W60258

Length: 339

February 11,

2000 15:48

Type: P

Check: 4443

MNQRNASMTV

IGAGSYGTAL

AITLARNGHE

VVLWGHDPEH

IATLERDRON

51

AAFLPDVPFP DTLHLESDLA TALAASRNIL VVVPSHVFGE VLRQIKPLMR

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New method for increasing production of 1,3-propane:diol - comprises fermentation of inexpensive carbon sources by microorganism expressing dehydratase, used, e.g. to prolong half-life of enzyme bisclosure; Page 70-71; 133pp; English.

The sequence is that of glycerol-3-phosphate dehydrogenase. It was used as part of a method of fermentative production of 1,3-propanediol (1,3-pd), using an organism comprising at least 1 gene encoding a dehydratase, is improved by inserting into the host a gene encoding protein x and culturing the transformant in presence of a carbon source (e.g. mono-, oligo-organism starting material for polyssters, polyurethanes and cyclic compounds. 1,3-pd can now be produced by a single recombinant organism from inexpensive carbon sources such as glucose (rather than costly glycerol or dihydroxyacetone), rapidly and without causing pollution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W54728 Length: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     residue(s) to antigen for increasing T cell response, useful in, e.g. vaccines against viral infection(s)

Disclosure; Page 34; 47pp; English.

The peptides W54559-W54609 are examples of peptides to which at least 1 (preferably 2) mannose can be attached to increase their uptake as antigens by antigen-presenting cells. Uptake of agonist mannosylated peptides will increase the T cell response, whereas uptake of antagonist peptides will increase the T cell response. Blocking binding of immunogenic autoantigens can be used in treatment of type I diabetes, rheumatoid arthritis, graft rejection etc., also to induce T-cell non-responsiveness. Vaccines containing mannosylated antigen are used to prevent or treat infections by, e.g. bacteria, viruses, fungi, helminths
                                                                                                                                                                                                                                                                                                               22-WAY-1998.
13-NOV-1997; U20873.
13-NOV-1996; US-030601.
(GEMY ) GENENCOR INT INC.
(GEMY ) DIAZ-TORTES M, Dunn-coleman NS, Trimbur NPI; 98-297944/26.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Klebsiella pneumoniae glycerol-3-phosphate dehydrogenase, glycerol-3-phosphate dehydrogenase; production;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            prevent or treat infections by, e. and parasites.
Sequence 24 AA;
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PDARLVWATK

GLEAETGRLL ODVAREALGD

QIPLAVISGP

TFAKELAAGL

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CC computing at least 1 gene encoding protein X and culturing Corpoly-saccharide or 1 gene encoding protest X and culturing CC or poly-saccharide or 1 gene encoding protein Sources by microvaganism CC recombination of inaxpensive carbon sources by microvaganism of fermentation of inexpensive carbon sources by microvaganism fermentation of inexpensive carbon sources by microvaganism protein used as part of a method of corpolation of 1,3-propandiol (1,3-pd), using an organism CC computising at least 1 gene encoding a dehydratase, is improved by CC computising at least 1 gene encoding protein X and culturing CC inserting into the host a gene encoding protein X and culturing CC or poly-saccharide or 1C substrate) convertible to 1,3-pd corpoly-saccharide or 1C substrate convertible to 1,3-pd corpoly-saccharide or 1C substrate) convertible to 1,3-pd corpolinate organism from inexpensive carbon sources such convertible convertible to 1,3-pd carbon corpolination cranism from inexpensive carbon sources such convertible convertible to 1,3-pd carbon convertible to 1,3-pd carbon corpolination cranism from inexpensive carbon sources such convertible convertible to 1,3-pd carbon convertible to 1,3-pd carbon convertible to 1,3-pd convertible to 1,3-
          ID AAC AC DT DE CS PN PPR PR PR PR PR
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13-NOV-1996; US-030601.
(GENY) GENENCOR INT
Chase MW, Disc
90 APR-1998.

30 APR-1998.

16-0CT-1997; GB-023173.

06-NOV-1996; GB-021681.

17-0CT-1996; GB-021681.
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                                                                                                                                    28-SEP-1998 (first entry)
Rainbow trout interleukin 1 beta.
interleukin 1 beta; ILI; beta; vaccine;
Oncorhynchus mykiss.
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lycerol-3-phosphate dehydrogenase;
3-propanediol; recombinant.
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                                                                                                                                                                                                                                              764 standard; Protein; 260
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!!AA_SEQUENCE 1.0 ID W44905 standard; AC W44905;

51 101 151 201 251

VTVDVPWSGS

SSKLCCVQYV

QAFEAKKDAS

AARASPSDVE WKSGKAGPGV LQPFTYELFV VAPVRVISHE GVTAEVFVKG AGELVAYSSA

SVKDVSQFAV
RAIKFAPIGL
TPRLCKVNGD

KCFSEFSVPL

LRPDEGVDLT

W57889 Length: 271

QSTHPCAAFH

AASRAISGGP IYVSDSVGQH

DFALLRRLAL PDGTVLRCEG

GGWSPEARRN YAVEARTLQI Type: P

Check: '5290

February 11, 2000 15:48

HALPTRDCLF

ADPLHDGRTV LKIWNVNRFA GVVGAFNCQG

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SCCCPTTTT
        !!AA_SEQUENCE 1.0
ID W57889 standard;
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pp 18-DEC-1999; 122417.

pr 18-DEC-1999; JP-338673.

pr (SUMO) SUMITOMO CHEM CO LTD.

pr (SUMO) SUMITOMO CHEM COME TAREFINOSE SYNTHETASE OF THE MICH COME OF THE MI
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WPI; 98-261498/23.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The sequence is that of rainbow trout interleukin 1 (IL-1) beta which can be used in the manufacture of a medicament for fish e.g. it can be used as a adjuvant in existing and developing vaccines. Sequence 260 AA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               existing and developing Claim 14; Fig 3; 46pp; The sequence is that o
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Raffinose synthetase; metabolism modification;
gastrointestinal flora; corn.
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EP-849359-A2.
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Loping fish vaccines
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PT Introducing genes into eukaryotic cells using peptide with two processors are specially as part of a viral glyco-protein, used in gene therapy to ensure precise targeting of PT glyco-protein, used in gene therapy to ensure precise targeting of PT vectors to particular cell types

PS Disclosure; Fig 36; 117pp; French.

CC A novel chimeric protein for gene transfer comprises, at the N-terminus, CC a protein binding a target molecule on the cell surface, and, at the CC terminus, a second protein recognising a secondary target molecule on the cell surface. The two domains are separated by a peptide comprising CC folds (beta or reverse turns) in the peptide sequence, the folds being CC regularly spread to form a polyproline beta-turn helix structure. This CC sequence is an example of a Pro-rich linking peptide. The pro-rich CC peptide acts by facilitating or inhibiting binding between the C-terminal CC protein does not interact with its target. The whole construct especially forms part of a (retro)viral envelope glycoprotein. The recombinant CC (retro)viral particles are used for in vitro or ax vivo transfer of CC nucleic acid into target cells, especially for gene therapy, e.g. in CC cystic fibrosis, myopathy, lysosomal disorders, cancer, viral (especially SC) Sequence 39 AA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "polyproline beta-turn helix" polypeptide Delta-PRO4-int.
"polyproline beta-turn helix" polypeptide Delta-PRO4-int.
Chimeric; polyproline; beta-turn helix; facilitation; inhibition;
retrovirus; envelope glycoprotein; gene therapy; cystic fibrosis;
myopathy; lysosomal disorder; cancer; human immunodeficiency virus.
Synthetic.
Synthetic.
Synthetic.
27-NOV-1997.
27-NOV-1997.
Introducing genes into eukaryotic cells using peptide with two receptor binding regions - especially as part of a viral glyco-protein, used in gene therapy to ensure precise targeting of vectors to particular cell types pisclosure; Fig 37; 117pp; French.

A novel chimeric protein for gene transfer comprises, at the N-terminus, a protein binding a target molecule on the cell surface, and, at the C-terminus, a second protein recognising a secondary target molecule on the cell surface. The two domains are separated by a peptide comprising at least 30% pro residues arranged so as to induce approximately 180 deg. Folds (beta or reverse turns) in the peptide sequence, the folds being regularly spread to form a polyproline beta-turn helix structure. This sequence is an example of a Pro-rich linking peptide. The pro-rich peptide acts by facilitating or inhibition occurs as long as the N-terminal protein and its receptor, and inhibition occurs as long as the N-terminal
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Synthetic.
W09744474-A2.
27-NOV-1997.
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20-MAY-1996; FR-006234.
(CNRS ) CENT NAT RECH SCI.
COSSET FL, Russell SJ, Valsesia
WFI; 98-018526/02.
N-FSDB; V30425.
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20-MAY-1996; FR-006234.
(CNRS ) CENT NAT RECH SCI.
COSSET FL, RUSSE11 SJ, Valsesia
WFI: 98-018526/02.
WPI: 98-018526/02.
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"polyproline beta-turn helix" polypeptide Delta-PRO4-beta.

Chimmeric; polyproline; beta-turn helix; facilitation; inhibition;

retrovirus; envelope glycoprotein; gene therapy; cystic fibrosis;

myopathy; lysosomal disorder; cancer; human immunodeficiency virus
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SEQUENCE 1.0 W44908 standard;

101

RQVLNVGPRV FOGATRGGRC

51

AAAETTGQAY

WKPTSSWDLI

SLKRGNTPWD

TGCSKVACGP CYDLSKVSNS Type: P

NPLVLEFTDA

29-SEP-1998 (first entry)

"polyproline beta-turn helix" polypeptide PRO-beta.

Chimeric; polyproline; beta-turn helix; facilitation; inhibition;

retrovirus; envelope glycoprotein; gene therapy; cystic fibrosis;

myopathy; lysosomal disorder; cancer; human immunodeficiency virus

Synthetic.

W09744474-A2.

27-NOV-1997.

W44907

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February

11,

2000 15:48

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              Introducing genes into eukaryotic cells using peptide with two proceeding genes into eukaryotic cells using peptide with two proceeding genes into eukaryotic cells using peptide with two proceeding genes in gene therapy to ensure precise targeting of proceeding in gene therapy to ensure precise targeting of proceding at gene transfer comprises, at the N-terminus, considered binding a target molecule on the cell surface, and, at the cell surface in the two domains are separated by a peptide comprising a target molecule on the cell surface. The two domains are separated by a peptide comprising cell teast 30% pro residues arranged so as to induce approximately 180 deg. Celds (beta or reverse turns) in the peptide sequence, the folds being regularly spread to form a polyproline beta-turn helix structure. This cell sequence is an example of a pro-rich linking peptide. The pro-rich cell sequence is an example of a pro-rich linking between the C-terminal protein does not interact with its target. The whole construct especially forms part of a (retro)viral envelope glycoprotein. The recombinant cell fibrosis, myopathy, lysosomal disorders, cancer, viral (especially sequence 127 AA:
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16-MAY-1997; F00870.4.
20-MAY-1996; FR-006234.
(CNRS ) CENT NAT RECH SCI.
(CNRS ) CENT NAT RECH SCI.
COSSet FL Russell SJ, Valsesia S;
WPI; 98-018525/02.
N-PSDB; V30426.
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"polyproline beta-turn helix" polypeptide Delta-PRO4-vrb.

Chimeric; polyproline; beta-turn helix; facilitation; inhibition; retrovirus; envelope glycoprotein; gene therapy; cystic fibrosis; myopathy; lysosomal disorder; cancer; human immunodeficiency virus
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Sequence

110 AA;

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receptor binding regions - especially as part of a viral plyco-protein, used in gene therapy to ensure precise targeting of provectors to particular cell types

Provectors to particular cell types

Provectors to particular cell types

Prove the chimeric protein for gene transfer comprises, at the N-terminus, a condition a target molecule on the cell surface, and, at the cerminus, a second protein recognising a secondary target molecule on the cell surface, and, at the cell surface. The two domains are separated by a peptide comprising the cell surface arranged so as to induce approximately 180 deg. CC at least 30% pro residues arranged so as to induce approximately 180 deg. CC regularly spread to form a polyproline beta-turn helix structure. This ceguence is an example of a pro-rich linking peptide. The pro-rich comprising peptide acts by facilitating or inhibiting binding between the C-terminal protein and its receptor, and inhibition occurs as long as the N-terminal composition of a (retro)viral envelope glycoprotein. The recombinant composition part of a (retro)viral envelope glycoprotein. The recombinant composite acid into target cells, especially for gene therapy, e.g. in cucleic acid into target cells, especially for gene therapy, e.g. in composite the content of composition of the composition of
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Process to particular cell types

Process to the cell surface, and the N-terminus, and the cell surface and the cell surface, and the cell surface and the cell surface, and the cell surface are separated by a peptide comprising cell the cell surface. The two domains are separated by a peptide comprising cell surface. The two domains are separated by a peptide comprising cell seast 30% processed the process to induce approximately 180 deg. Cell the two the cell surface, the folds being cell surface to form a polyproline beta-turn helix structure. This cell sequence is an example of a pro-rich linking peptide. The pro-rich cell sequence is an example of a pro-rich linking peptide. The pro-rich cell peptide acts by facilitating or inhibiting binding between the C-terminal cell protein does not interact with its target. The whole construct especially forms part of a (retro)viral envelope glycoprotein. The recombinant cell construct and into target cells, especially for gene therapy, e.g. in custoff brossis, myopathy, lysosomal disorders, cancer, viral (especially construct construct cells) into the first cells.
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27-NOV-1997;
16-MAY-1997; F00870.
20-MAY-1996; FR-006234.
(CNRS) CEWT NAT RECH SCI.
(CNRS) CEWT NAT RECH SCI.
COSSET FT., RUSSELI SJ. VALSESIA S
WPI; 98-018526/02.
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N-PSDB; V30427.
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20-MAY-1996; FR-006234.
(CNRS) CENT NAT RECH S
Cosset FL, Russell SJ,
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IID AA ID AC DIT DE DE KW KW KW KW KW KW KW FN OS PD PF

Synthetic. WO9744474-A2.

F00870. FR-006234

SEQUENCE 1.0 w44879 standard;

Protein;

104

8

151 101

51 _

RQVLNVGPRV FQGATRGGRC

PIGPNPVLPD NPLVLEFTDA WKPTSSWDLI

QRLPSSPIEI VPAPQPPSPL

GKKANWDGPK SLKRGNTPWD

SWGLRLYRIG TGCSKVACGP

TDPITMESLT CYDLSKVSNS

VPQPPPA

AAAETTGQAY

W44879;
29-SEP-1998 (first entry)
29-SEP-1998 (first entry)
"polyproline beta-turn helix" polypeptide PRO+(4070A).
"polyproline beta-turn helix; facilitation; inhibition; Chimeric; polyproline; beta-turn helix; facilitation; tibrosis; retrovirus; envelope glycoprotein; gene therapy; cystic fibrosis; cancer; human immunodeficiency virus; cancer; h

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Introducing genes into eukaryotic cells using peptide with two preceptor binding regions - especially as part of a viral preceptor binding regions - especially as part of a viral preceptor binding regions - especially as part of a viral preceptor binding regions - especially as part of a viral preceptor binding at a particular cell types prench.

CC aprotein binding a target molecule on the cell surface, and, at the ccl aprotein binding a target molecule on the cell surface and, at the ccl extensions, a second protein recognising a secondary target molecule on the cell surface. The two domains are separated by apeptide comprising the cell surface. The two domains are separated by a peptide comprising consequence is an example of a pro-rich linking peptide. The pro-rich generated acts by facilitating or inhibiting binding between the C-terminal protein and its receptor, and inhibiting binding between the C-terminal protein does not interact with its target. The whole construct especially constituted acts by facilitating or inhibiting binding between the construct especially construct approach of a (retro)viral envelope glycoprotein. The recombinant construct approach of a construct especially for gene therapy, e.g. in construct of a construct especially for gene therapy, e.g. in the construct of the construct especially for gene therapy, e.g. in construct of the construct of the
W44910 Length: 167
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W09744474-A2.
27-NOV-1997. F00870.
16-MAY-1997. F00870.
20-MAY-1996. FR-006234.
(CORS ) CENT NAT RECH SCI.
COSSET FL, RUSSELL SJ, Valsesia S
WPF, 98-018526/02.
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W44910 standard;
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Chimeric; polyprolino
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Chimeric; polyproline; beta-turn helix; facilitation; inhibition;
retrovirus; envelope glycoprotein; gene therapp; cystic fibrosis;
myopathy; lysosomal disorder; cancer; human immunodeficiency virus.
                                                         Sequence
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                                                         167 AA;
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      February 11, 2000 15:48
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      Type: P
            Check: 1990
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Introducing genes into eukaryotic cells using peptide with two receptor binding regions - especially as part of a viral glyco-protein, used in gene therapy to ensure precise targeting of precise to particular cell types

PT vectors to particular cell types

S Disclosure; Fig 13; 117pp; French.

CC A novel chimeric protein for gene transfer comprises, at the N-terminus, cc cell surface are comparted for gene transfer comprises, at the N-terminus, cc cell surface. The two domains are separated by a peptide comprising cat least 30% pro residues arranged so as to induce approximately 180 deg. Cc terminus spread to form a polyproline beta-turn helix structure. This sequence is an example of a Pro-rich linking peptide. The pro-rich peptide acts by facilitating or inhibiting binding between the C-terminal protein and its receptor, and inhibition occurs as long as the N-terminal protein does not interact with its target. The whole construct especially forms part of a (retro)viral envelope glycoprotein. The recombinant curleic acid into target cells, especially for gene therapy, e.g. in human immunodeficiency virus) infections, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PT Introducing genes into eukaryotic cells using peptide with two receptor binding regions - especially as part of a viral proceeding yence in gene therapy to ensure precise targeting of plyco-protein, used in gene therapy to ensure precise targeting of process to particular cell types

PT vectors to particular cell types

S Disclosure; Fig 10; 117pp; French.

CC A novel chimeric protein for gene transfer comprises, at the N-terminus, cell surface to the cell surface. The two domains are separated by a peptide comprising cell telest 30% pro residues arranged so as to induce approximately 180 deg.

CC terminus, a second protein recognising a secondary target molecule on the cell surface. The two domains are separated by a peptide comprising cell telest 30% pro residues arranged so as to induce approximately 180 deg.

CC terminus as receptors as topinde sequence, the folds being cell telest so a example of a province beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproline beta turn helix structure. This cegunary spread to form a polyproli
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20-MAY-1996; FR-006234.
(CNRS ) CENT NAT RECH SCI.
COSSEC FL, RUSSELL SJ, Valsesia S;
WPI; 98-018526/02.
N-PSDB; V30401.
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Cosset FL, Russell
WPI; 98-018526/02.
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Russell SJ, Valsesia
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VSQGTPMAPA PYPQSPFPPN

PYGQPQVFPG

QDPDSPQHGN

YYEEGPPSYY

DNQDFPATNW

PTPKGATHRV

PTPKGATHRG

VLSCCGDFRR

KHPWNLVALS

LSVTLSTVSV

ASLGALLFTC

MOTRYDFTSC

MGVLLVSMVV

MKRVFWCLGT TILPPTLDIR GGPSHPCPPM LSLPTLGPLT

HSPLSSPPPT

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IIAA_SEQUENCE 1.0

AC W625612 standard; Protein; 369 AA C W625612

DIE Allegard (first entry)
DE Human glutamate-binding protein KW Human; glutamate-binding protein KW treatment; prevention; diagnosis kW signalling; glutamate; ischaemia kW schizophrenia; neurodegeneration KW lathyrism.

OS Homo sapiens. Location/Qualifi FT Region 7/00te= "transmem FT Region 201.290". 320

FT Region 207.320

FT Nov-1997; U20860.

FT No-PSDB; U3-749289.

PA 14-NOV-1997; U20860.

FT No-PSDB; U3-749289.

FT No-PSDB; U3-749289.

FT No-PSDB; U3-749289.

FT No-PSDB; U3-749289.

FT Nov-1997; U20860.

FT PORT Nov-1997; U20860.

FT PORT Nov-1997; U20860.

FT Region 207.320

FT PORT Nov-1997; U20860.

FT Region 207.320

FT PORT Nov-1997; U20860.

FT Region 207.320

FT Reg
                                     W62612
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Human; glutamate-binding protein; HGLUBP; Incyte clone 386116;

treatment, prevention; diagnosis; central nervous system disorder;

signalling; glutamate; ischaemia; hypoglycaemia; epilepsy; convuls

schizophrenia; neurodegeneration; amyotrophic lateral sclerosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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         Length: 369
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   February 11, 2000
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178. .21
note= "transmembrane"
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e= "transmembrane
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            protein - useful for treatment,
central nervous system diseases,
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   15:49
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Type: P
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!!AA_SEQUENCE 1.0

standard; Protein;

387

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From thathive production of 1,3-propanediol - by single organism retreatable production of 1,3-propanediol - by single organism retreatable comprising specific genes, and capable of using retreatable sources carbon sources.

Claim 13; Page 81-82; 95pp; English.

This is the 1,3 propanediol oxidoreductase encoded by the dharmous specific gene (see va2015) of Klebstella pneumoniae AFCC 25955.

Cenzyme catalyses the reduction of 3-hydroxypropionaldehyde to 1,3-propanediol (1). A claimed method for production of (1) comprises culturing a microorganism transformed with a cassette containing at least 1 of the genes (see v42012-21) for glycerol-3-phosphate dehydrogenase, glycerol-3-phosphatase, glycerol-3-phosphatase, glycerol-3-phosphatase, and 1,3-propanediol oxidoreductase (see w30676-85). Any of these genes not present on the cassette must be present endogenously. The enzyme sequences may include substitutions, deletions and additions provided activity is not altered. A single recombinant organism can now be used for production of (1) from inexpensive C-sources (contrast use of a starting material for polyesters, polyurethanes and cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             W30679
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Nagarajan V, Nair RV, Nakamura
Trimbur DE, Whited GM;
WPI; 98-297942/26.
N-PSDB; V42015 and V42022.
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W69232 stand
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13-NOV-1996; US-030601
(DUPO ) DU PONT DE NEM
(GEMY ) GENENCOR INT I
                                                                       Fc receptor-like protein; phagocytosis inducer; rheumatoid arthritis; immune complex related disease; systemic lupus erythematosus; allergy haemolytic anaemia; thrombocytopaemia; anaphylaxis; cancer; lymphoma; leukaemia; infection; immunomodulator; viral entry inhibitor; therapy
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              Peptide
                                          iomo sapiens.
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,3-Propanediol oxidoreductase,
,3-Propanediol oxidoreductase;
[Lebsiella pneumoniae ATCC 25955]
                                                                                                                                                  0-OCT-1998 (first entry)
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                                                                                                                                  protein
                                                                                                                                                                                standard; Protein;
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                                                                                                                                    sequence.
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                            Location/Qualifiers
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Wakamura CE,
"signal peptide
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Payne MS, Picataggio
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, and capable of using
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                                                                                       is; allergy;
lymphoma;
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DR N-PSDB; V44825.

PT Nucleic acid encoding Fc receptor-like polypeptides or their Nucleic acid encoding Fc receptor-like polypeptides or their provides. PT Nucleic acid encoding Fc receptor transformed cells and antibodies, provided for treating and diagnosing diseases of the haematopoietic provening for treating and diagnosing diseases of the haematopoietic provening for the Fc receptor-like II protein (FCR-II) of the Claim 23; Fig 2A; 141pp; English.

CC This sequence is the Fc receptor-like II protein (FCR-II) of the complex related diseases (e.g. rheumatoid surthritis, systemic lupus complex related diseases (e.g. rheumatoid arthritis, systemic lupus complex related diseases (e.g. rheumatoid arthritis, systemic lupus complex related diseases (e.g. rheumatoid arthritis, systemic lupus complex related of sease (e.g. rheumatoid arthritis, systemic lupus complex related of sease (e.g. rheumatoid arthritis, systemic lupus complex related of inhibitors of viral (e.g. human immune deficiency or immunomodulators and inhibitors of viral (e.g. human immune deficiency or dengue viruses) entry into cells. The proteins may also be used to screen (ab), and for identification of particular cells or raising antibodies (ab), and for identification of particular cells or tissues. The Ab can be used therapeutically as antagonists, for raising antibodies (affinity purification of the proteins. The DNA and its fragments are cuseful as hybridisation (chromosome mapping) and diagnostically to measure (c. g. semmence 163 AA.
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23-JUL-1998.
20-JAN-1998; U01184.
18-JUN-1997; US-034205.
21-JAN-1997; US-034205.
(HUMA-) HUMAN GENOME SCI IN
GENTZ RL, MUTPHY M, NI J, C
WPI; 98-414105/35.
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Sequence
    263 AA;
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/note=
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Olsen HS, Ruben
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W69232 Length: 263 February 11, 2000 15:49 Type: P Check: 3029

 \vdash LRCRAPQPAW MALVLILQLL TLWPLCHTDI TPSVPPASYH PKPWLGAQPA REGLEKPGEI APLLFRDVSS ELAEFFLEEV TPAQGGSYRC TVVTPGVNVT

51

AGRLRNMSFV CYRRPDWGPG LYREGVAAPL VWSQPSDVLE LLVTEELPRP QYRHSAQPWA DFTLLGARAP SLVALPGPVV GPGANVSLRC GTYSCYYHTP

SAPYVLSQRS EVLVISWEDS GSSDYTRGNL VRLGLAGLVL ISLGALVTFD

201 151 101

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31-OCT-1997; AU0729.
14-FEB-1997; AU-005117.
14-FEB-1997; AU-005117.
01-NOV-1996; AU-003384.
(HALL-) HALL INST MEDICAL RES WALTER & ELIZA.
Alexander WS, Hilton DJ, Metcalf D, Nicholson SE,
Alexander WS, Hilton DJ, Metcalf D, Wicholson SE,
Nicola NA, Richardson RT, Starr R, Viney EM, Willson
                                                                                                                                                                                         27-OCT-1998 (first entry)
Rattus norvegicus SOCS1 protein.
SOCS; suppressor of cytokine signalling; PCR primer;
autoimmune disease; diagnosis; cancer; treatment;
cytokine mediated cellular responsiveness; hyperimmunity;
immunosuppression; allergies; hypertension.
Rattus norvegicus.
W09820023-A1.
                                                                                                                                                                                                                                                                                                                                      SEQUENCE 1.0
W62617 stand
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                                                98-286854/25.
)B; V38663
                                                                                                                                                                                                                                                                                                                                        standard; Protein;
cytokine signalling proteins - useful to treat
ry or abnormality involving cytokine mediated co
e.g. hyperimmunity, immunosuppression, allerg
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   cellular
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W62622
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                                                                                        hypertension
Claim 13; Page 149-150; 325pp; English.
Claim 13; Page 149-150; 325pp; English.
The sequence is that of a suppressor of cytokine signalling protein (SOCS). SOCS can be used to screen for naturally occurring antibodies to SOCS, which may occur, e.g. in some autoimmune diseases. Alternatively, specific antibodies can be used to screen for SOCS, which is useful as a knowledge of SOCS levels may be important for the diagnosis of certain cancers. Soluble SOCS polypeptides can be used to treat disease, injury or abnormality involving cytokine mediated cellular responsiveness, e.g. hyperimmunity, immunosuppression, allergies and hypertension. Sequence 130 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    hypertension
Claim 13; Pag
The sequence
protein (SOCS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 13; Page 119-120; 325pp; English.
The sequence is that of a suppressor of cytokine signalling protein (SOCS). SOCS can be used to screen for naturally occurring antibodies to SOCS, which may occur, e.g. in some autoim diseases. Alternatively, specific antibodies can be used to screen for SOCS, which is useful as a knowledge of SOCS levels may be important for the diagnosis of cerem. Sociable SOCS polypeptides can be used to treat disease, injury or abnormality involving cytokine mediated cellular responsiveness, e.g. hyperimmunity, immunosuppression, allergies and hypertension.
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31-OCT-1997; AU0729.
14-FEB-1997; AU-005117.
10-NOV-1996; AU-005317.
01-NOV-1996; AU-003384.
(HALL-) HALL INST MEDICAL RES WALTER & ELIZA.
Alexander WS, Hilton DJ, Metcalf D, Nicholson SE,
Alexander WS, Hilton DJ, Starr R, Viney EM, Willson
                                                                                                                                                                                                                                                                                                                                                           Suppressor
disease, in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mus musculus SOCS8 protein.
SOCS; suppressor of cytokine signalling; PCR primer;
autoimmune disease; diagnosis; cancer; treatment;
cytokine mediated cellular responsiveness; hyperimmunity;
immunosuppression; allergies; hypertension.
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                                                                                                                                                                                                                                                                                                                                   Suppressor of cytokine signalling proteins - useful to treat disease, injury or abnormality involving cytokine mediated celluresponsiveness e.g. hyperimmunity, immunosuppression, allergies
                                                                                                                                                                                                                                                                                                                                                                                                WPI; 98-286854/25.
N-PSDB; V38675.
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                                                                       Length: 130
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RVVQLPEEAK
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GLVPPEILQK YHGFYSSLFA
                                    CWLPVTSAVN
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                                                                         February 11,
                                    PORMLRPPPT AVENCANCCC
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LVRQPRSLQH
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LCRCALRSHI
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BAB

!!AA_SEQUENCE 1.0
ID W62623 standard; Protein;

suppressor

of cytokine signalling

EGCLPHALPR LPLPPRMLRF LQLDFEDLLY

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W62623
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Claim 13; Page 158-159; 325pp; English.
The sequence is that of a suppressor of cytokine signalling protein (SOCS). SOCS can be used to screen for naturally occurring antibodies to SOCS, which may occur, e.g. in some autoim diseases. Alternatively, specific antibodies can be used to screen for SOCS, which is useful as a knowledge of SOCS levels may be important for the diagnosis of certain cancers. Soluble sock polypeptides can be used to treat disease, injury or abnormality involving cytokine mediated cellular responsiveness, e.g. hyperimmunity, immunosuppression, allergies and hypertension.

sequence 207 AA;
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31-OCT-1997; AU0729.
14-FEB-1997; AU-005117.
11-NOV-1996; AU-003384.
(HALL-) HALL INST MEDICAL RES WALTER & ELIZA.
Alexander WS, Hilton DJ, Metcalf D, Nicholson SE,
Nicola NA, Richardson RT, Starr R, Viney EM, Willson
WPI: 98-286554/25.
hypertension hypertension 13; Page 164; 325pp; English. Claim 13; Page 164; 325pp; English.
                                                            Suppressor of cytokine signalling proteins - useful to treat disease, injury or abnormality involving cytokine mediated celluresponsiveness e.g. hyperimmunity, immunosuppression, allergies
                                                                                                                                  14-MAY-1998.
31-OCT-1997; AU0729.
14-FEB-1997; AU-005117.
14-FEB-1997; AU-005117.
01-NOV-1996; AU-003384.
(HALL-) HALL INST MEDICAL RES WALTER & ELIZA.
Alexander WS, Hilton DJ, Metcalf D, Nicholson SE,
Nicola NA, Richardson RT, Starr R, Viney EM, Willson
WPI; 98-286854/25.
                                                                                                                                                                                                                                                                                                                                              27-0CT-1998 (first entry)
27-0CT-1998 (first entry)
Mus musculus SOCS13 protein.
SOCS; suppressor of cytokine signalling; PCR primer;
autoimmune disease; diagnosis; cancer; treatment;
cytokine mediated cellular responsiveness; hyperimmunity;
immunosuppression; allergies; hypertension.
Mus musculus.
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                                                                                                                    WPI; 98-286854/25.
N-PSDB; V38685.
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SOCS; suppressor of cytckine signalling; PCR primer;
autoimmune disease; diagnosis; cancer; treatment;
cytckine mediated cellular responsiveness; hyperimmu
immunosuppression; allergies; hypertension.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         standard; Protein; 134
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    ACC DT DE KW
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W62613 standa
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Claim 13; Page 110; 325pp; English
The sequence is that of a suppressor of cytokine signalling
protein (SOC5). SOCS can be used to screen for naturally
protein (SOC5). SOCS can be used to diseases. Alternatively, specific antibodies can be used to
diseases. Alternatively, specific antibodies can be used to
screen for SOCS, which is useful as a knowledge of SOCS lovels
screen for SOCS, which is useful as a knowledge of SOCS lovels
may be important for the diagnosis of certain cancers. Soluble
may be important for the diagnosis of certain cancers. Soluble
SOCS polypeptides can be used to treat disease, injury or
SOCS polypeptides can be used to treat disease, injury or
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Mus musculus SOCS1 protein
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31-OCT-1997; AU0729.
14-FEB-1997; AU-005117.
10-NOV-1996; AU-003384.
(1ALL-) HALL INST MEDICAL RES WALTER & ELIZA.
Alexander WS, Hilton DJ, Metcalf D, Nicholson Alexander WS, Hilton BJ, Metcalf D, Nicholson RI, Starr R, Viney EM, W MPI; 98-286584/25.
N-PSDB; V38659.
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SOCS; suppressor of cytokine signalling; PCR primer;
autoimmune disease; diagnosis; cancer; treatment;
cytokine mediated cellular responsiveness; hyperimmunity;
immunosuppression; allergies; hypertension.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            suppressor of cytokine signalling proteins - useful to treat disease, injury or abnormality involving cytokine mediated cell responsiveness e.g. hyperimmunity, immunosuppression, allergies
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27-OCT-1998 (first entry)
Mus musculus SOCS2 protein
SOCS; suppressor of cytokir
autoimmune disease; diagno
                                                                                                                                                                                                                                                                                                                                                                              151
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                                                                                                                                                                                       W62614 standard;
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                                                                                                                                                                                           Protein; 198
             cytokine signalling; PCR primer; diagnosis; cancer; treatment;
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be used to
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IIAA_SEQUENCE 1.0

WESS15;

DT WESS15;

WESS15;

WESS15;

WESS25;

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Claim 13; Page 112; 325pp; English.

Claim 13; Page 112; 325pp; English.

Claim 13; Page 112; 325pp; English.

The sequence is that of a suppressor of cytokine signalling

The sequence is that of a suppressor of cytokine signalling

protein (SOCS), SOCS can be used to screen for naturally

protein (SOCS), SOCS can be used to diseases. Alternatively, specific antibodies can be used to

diseases. Alternatively, specific antibodies can be used to

screen for SOCS, which is useful as a knowledge of SOCS levels

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     suppressor of cytokine signalling proteins - useful to treat disease, injury or abnormality involving cytokine mediated cellular responsiveness e.g. hyperimmunity, immunosuppression, allergies and hyperimmunity.
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N-PSDB; V38660.
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(HALL-) HALL INST MEDICAL RES WALTER & ELIZA.
Alexander WS, Hilton DJ, Metcalf D, Nicholson
Nicola NA, Richardson RT, Starr R, Viney EM, V
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Mus musculus SOCS3 protein.
SOCS; suppressor of cytokine signalling; PCR primer;
succismmune disease; diagnosis; cancer; treatment;
cytokine mediated cellular responsiveness; hyperimmunity;
immunosuppression; allergies; hypertension.
hypertension

Claim 13: Page 114-115; 325pp; English.

Claim 13: Page 114-115; 325pp; English.

Claim 13: Page 114-115; 325pp; English.

The sequence is that of a suppressor of cytokine signalling

The sequence is that of a suppressor of cytokine signalling

The sequence is that of a suppressor of cytokine signalling

The sequence is that of a suppressor occur, e.g. in some autoims

occurring antibodies to socs, which may occur, e.g. in some autoims

other sequence is some autoims of series of socs levels

screen for SOCS, which is useful as a knowledge of SOCS levels

screen for SOCS, which is useful as a knowledge of SOCS levels

screen for SOCS, which is useful as a knowledge of SOCS levels

screen for SOCS, which is useful as a knowledge of SOCS levels

screen for SOCS polypeptides can be used to treat disease, injury or

SOCS polypeptides can be used to treat disease, injury or

SOCS polypeptides can be used to treat disease, injury or

social interpretation of the social interpreta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    suppressor of cytokine signalling proteins - useful to treat disease, injury or abnormality involving cytokine mediated celluresposiveness e.g. hyperimmunity, immunosuppression, allergies
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31-QCT-1997; AU0729.
14-FEB-1997; AU-005117.
10-NOV-1996; AU-003184.
(HALL-) HALI INST MEDICAL RES WALTER & ELIZA.
(HALL-) HALI INST MEDICAL RES WALTER & ELIZA.
Nicola NA, Richardson RT, Starr R, Viney EM, W
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Willson TA;
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The sequence is that of a suppressor of cytokine signalling protein (SOCS). SOCS can be used to screen for naturally occurring antibodies to SOCS, which may occur, e.g. in some autoimmune diseases. Alternatively, specific antibodies can be used to screen for SOCS, which is useful as a knowledge of SOCS levels may be important for the diagnosis of certain cancers. Soluble SOCS polypeptides can be used to treat disease, injury or abnormality involving cytokine mediated cellular responsiveness, e.g. hyperimmunity, immunosuppression, allergies and hypertension.
                                                                                                          27-OCT-1998 (first entry)
Clone HTPEF86 of TM4SF superfamily.
Human: receptor; immune disorder; canc
juvenile rheumatoid arthritis; Graves
                                                                                                                                                                                                                                                        SEQUENCE 1.0
W61619 standard; Protein;
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14-FEB-1997; AU-005117.
11-FEB-1997; AU-005117.
01-NOV-1996; AU-003384.
(HALL-) HALL INST MEDICAL RES WALTER & ELIZA.
(HALL-) HALL INST MEDICAL RES WALTER & ELIZA.
Alexander WS, Hilton DJ, Metcalf D, Nicholson
Nicola NA, Richardson RT, Starr R, Viney EM, W
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SOCS; suppressor of cytokine signalling; PCR primer;
autoimmune disease; diagnosis; cancer; treatment;
cytokine mediated cellular responsiveness; hyperimmunity;
immunosuppression; allergies; hypertension.
                                                      Homo sapiens.
W09831799-A2.
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23-JUL-1998.
21-JAN-1998; U00959
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W65460

Length:

375

February 11,

2000 15:49

Type: P

Check:

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Transgenic animal growth differentiation factor-11 is inhibited by insertion of transgene, also use of GDF-11 inhibitors for treating muscular wasting, neuromuscular disease, obesity

Reample 3; Page 55-56; 89pp; English.

This 1s the amino acid sequence of human growth differentiation can be used human growth differentiation factor-11 to the newly identified human growth differentiation factor-11 compared to the newly identified human growth differentiation factor-11 sequences protectial N-linked glycosylation signals and putative reveals potential N-linked glycosylation signals and putative reveals potential N-linked glycosylation signals and putative protection is sequences are related not only in the C-terminal region following the putative cleavage site (90% amino acid sequence identity) but also in the pro-region of the molecules (45% amino acid sequence identity. Claimed transgenic animals in which GDF-11 production is reduced produce higher than normal levels of muscle and are useful in the food industry. GDF-11 polypeptides, polynucleotides and antibodies can be used to modulate GDF-11 activity or gene expression for treatment of cell proliferative disorders involving sequence 375 AA;
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13-AUG-1998; U02310.
06-FEB-1997; US-795671.
(UYJO) UNIV JOHNS HOPKL
Lee S, McPherron AC;
WFI; 98-447217/38.
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WPISDB; V48114.

N-PSDB; V48114.

New isolated poly:nucleotide(s) and encoded receptor poly:peptide(s)

- used to develop products for diagnosing or treating e.g. immune

- used to develop products for diagnosing or immuno-compromised disease
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Clone HTPEF66 is a member of the TM4SF receptor superfamily. The products generated using the receptor can be used for treating abnorm conditions related to both an excess of and insufficient amounts of receptor activity. They can be used in the treatment of e.g. immune disorders, cancers, blood disorders, juvenile rheumatoid arthritis, Graves disease or immunocompromised disease states. The products can sloo be used for detection and diagnosis.
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Human growth differentiation factor-8.
Growth differentiation factor-8; GDF-8;
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21-JAN-1997; US-034204.
(HUMA-) HUMAN GENOME SC
Gentz RL, Ni J, Rosen C
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263. .266
/note= "RXXR proteolyt1
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Mycobacterium tuberculosis antigen RDIF5.
Tuberculosis; infection; diagnosis; antigen; RDIF5.
Mycobacterium tuberculosis strain Erdman.
Mycobacterium tuberculosis antigen RDIF5.
Mycobacterium tuberculosis strain Erdman.
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                                    Streptococcus I
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W38580 star
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Campos-Neto A, Dillon DC, Houghton R, Lodes MJ,
Reed SG, Skelky YAW, Twardzik DR, Vedvick TS;
WPI, 98-251292/22.
                                                                                                                                                                                                                              W38580;
06-NOV-1998 (first entry)
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                                    pneumoniae protein of unknown function. pneumoniae protein; genetic immunisation; an response; inoculation; antibody production; response; antimicrobial compound; bacterial matrix protein; protein-mediated cell invasi
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                                                                                 antagonist; on; inhibitor; lal adhesion;
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Claim 12: Pages 351-352; 483pp: English.

CC This sequence represents a Streptococcus pneumoniae protein of this sequence represents a Streptococcus pneumoniae of the invention. CC Inknown function, and is encoded by a DNA sequence of the invention. CC The DNA sequences were isolated from Streptococcus pneumoniae strain cc 1010993 (NCIMB 40794). The Streptococcus pneumoniae proteins of the cinhibit or activate the activity of the proteins. Antagonists can be used to identify compounds which interact with and cc inhibit or activate the activity of the proteins. Antagonists can be used to treat diseases caused by s. pneumoniae proteins, through genetic used to treat diseases caused by s. pneumoniae proteins or delivery. CC of the encoding nuclei acids in a vector adequate to produce antibody and/or T cell immune responses to protect the animal from disease. The CC capable of inhibiting their bloactivity. In particular the proteins of the invention can be used to prevent adhesion of bacteria to mammalian CC extracellular matrix proteins on in-dwelling devices or in wounds, to block protein-mediated mammalian cell invasion, and to block the normal CC implantation of in-dwelling devices or in wounds. Sc sequence 174 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   W38580
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14-MAY-1997; U07950.

14-MAY-1997; U07950.

14-MAY-1996; US-017670.

(SMIK ) SMITHKLINE BEECHAM CORP.

(SMIK ) SMITHKLINE BEECHAM PLC.

(SMIK ) SMITHKLINE BECHAM PLC.
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Key
Location/Qualifiers
Misc_difference 1. .174
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        WPI; 98-399137/34.

N-PSDB; V42701.

STAT function regulatory protein - used in screening candi substances for cytokine regulatory activity

Claim 2; Pages 39-41; 60pp; Japanese.

The present sequence represents a protein (called SIIS-1, inhibitor of STAT function 1) which regulates STAT protein the JAK/STAT signal transmission system in mammalian cells
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                                                                                                                                                                                                                                                                       A STAT function regulatory protein designated SIIS-1.

SIIS-1, STAT-induced inhibitor; STAT function;

JAK/STAT signal transmission system; STAT3; STAT6; inhibit;

tyrosine phosphorylation; gpi30; cytokine-regulating protein;

screen; cytokine regulatory; inhibitory activity.
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TAT protein lian cells

, STAT-induced in function in lanction in lanction in lanction.

candidate

is induced by STAT3 or STAT6. It inhibits tyrosine phosphorylation STAT3 and of gpl30. The SIIS-1 protein sequence contains an SH2 don and is related to the cytokine-regulating protein CIS. SIIS-1, or transformant cells expressing it, may be used to screen candidate substances for cytokine regulatory or inhibitory activity. sequence 212 AA;

on of

PXIFIGTALP LAGTVATGAI

HFTANEVIPI

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New nucleic acid sequences from Staphylococcus aureus WCHU29 - useful in vaccines and for treatment of bacterial infections of e.g. respiratory tract and central nervous system claim 11; Page 317-318; 390pp; English.

Chaim 12; Page 317-318; 390pp; English.

Chaim 13; Page 317-318; 390pp; English.

Chaim 14; Page 317-318; 390pp; English.

Chaim 15; Page 317-318; 390pp; English.

Chaim 16; Page 317-318; 
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W77666 standa
W77666;
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(SMIK ) SI
Black MT,
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13-MAY-1998.
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Staphylococcus aureus protein; immune response induction; eye infection;
antibody production; T-cell immune response; gastrointestinal infection;
respiratory infection; inhibitor; bacterial infection; cardiac infection
central nervous system; kidney infection; urinary tract infection;
antimicrobial compound identification; broad spectrum antibiotic;
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identify antimicrobial conspecially useful in the Sequence 100 AA;
                                                                                                                                                                                                                                                                                                                                                                                    Ward JM;
WPI; 98-252940/23.
N-PSDB; V53459.
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24-SEP-1996; US-027032
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Lonetto MA, Nicholas RO,
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SMITHKLINE BEECHAM PLC.
T, Burnham MKR, Hodgson JE,
MA, Nicholas RO, Pratt JM,
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42
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Reichard RW, Rosenberg
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W77666

Length:

100

February 11,

2000

15:49

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Check:

PR PR PR

A Staphylococcus aureus protein.
Alsaligenes eutrophus; catton efflux system czcD protein; treatment; prevention; bacterial infection; Helicobacter pylori; vaccine.

SEQUENCE 1.0 W79302 standard; Protein;

67

AΑ

24-NOV-1998 (first entry)

101

51

GLEPLTADNY

IQEELTTALA LEDKDLVLFV ADTLEVANAT LGALRGRIAK ELGLIDNDKF

SRKDIDKMTE VAKQYGAKGL AWVKVVDGEL

P Check: 3773
ALAVKAIVVN
NGPVAKFLTG

MLLQDLTEVV

KGVDFKVFSE A

MKYDDAMALY GSDKPDTRFD

04-JUN-1998. 24-NOV-1997; 25-NOV-1996;

Staphylococcus WO9823738-A2. W38716 Length: 161

February 11,

2000 15:49

Add Mark and Supply Sup

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W79302 Length: 67 February 11,
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06-AUG-1998.
05-FEB-1998; U02479.
23-MAY-1997; US-862445.
05-FEB-1997; US-847910.
28-APR-1997; US-847910.
28-APR-1997; US-847910.
28-DES, MCPHORION AC;
WPI; 98-437444/37.
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Rat growth differentiation factor-8.
Rat growth differentiation factor-8; GDF-8; rat; transgenic animal; growth differentiation factor-beta; muscle; meat; inhibitor; obesity; transforming growth factor-beta; muscle; cachexia; AIDS; cancer; neuromuscular disease; muscular dystrophy; cachexia; AIDS; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Transgenic animals with gene for growth differentiation factor-8 disrupted - have increased muscle and reduced cholesterol contents, also use of GDF-8 inhibitors for treating cancer, obesity,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cleavage_site
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264. .267
268. .376
/label= Mat_protein
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8888888888888
        W69889 Length: 376
Example 9; Fig 14f; 125pp; English.

Example 9; Fig 14f; 125pp; English.

This is the amino acid sequence of the transforming growth factor.

Generally that appears to relate to various cell; the person of the transforming growth factor.

Deta superfamily that appears to relate to various cell; nerve deta superfamily those appears to relate to various cell; nerve and adipose tissue. The sequence was deduced from a cDNA clone and adipose tissue. The sequence was deduced from a cDNA clone can be sequence with the sequence with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and for poultry reduced cholesterol contents. Method (3) is used to treat muscle wasting or neuromuscular diseases, muscular atrophy and aging, particularly muscular dystrophy, spinal cord or traumatic injuries, congestive or obstructive lung disease, AIDS and cachexia. Method (4) is used to treat cancer of muscle, and connective tissue and bone, or obesity. Also (not claimed) GDF-8 can be used to maintain myoblasts intended for transplanting or to improve efficiency of fusion. Ab can be used to detect and quantify GDF-8 (particularly in muscle, for diagnosis or monitoring), also for immunotherapy and in vivo imaging.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      05-AUG-1998

05-FEB-1998; UO2479.

23-MAY-1997; US-862445.

05-FEB-1997; US-95071.

28-APR-1997; US-647910.

(UYJO) UNIV JOHNS HOPKI

Lee S, McPhetrion AC;

WPI: 98-437444/37.
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Sus scrofa.
Key
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07-DEC-1998 (first entry)
Pig growth differentiation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard; Protein;
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263..266
267..375
/label= Mat_protein
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; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                   contents,
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, 888888888888888

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r neuromiscular disease

Stample 9; Fig 14f; 125pp; English.

This is the amino acid sequence of sheep growth differentiation

factor-8 (GDF-8), a novel member of the transforming growth factor-

beta superfamily that appears to relate to various cell

proliferative disorders, especially those involving muscle, nerve

and adipose tissue. The sequence was deduced from a cDNA clone

(see V45823) isolated from a skeletal muscle cDNA library. The

invention provides novel mammalian and avian GDF-8 proteins (see

W69883-92). A transgenic non-human animal is claimed in which

GDF-8 expression is disrupted or interfered with. Also claimed

are: (1) chicken or turkey eggs or meat, beef, milk, pork and lamb

from these animals; (2) method for increasing muscle mass in

animals by administering an antibody (Ab) that binds to GDF-8; (3)
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06-AUG-1998. U02479.
05-FEB-1998; US-862445.
05-FEB-1997; US-959071.
28-APR-1997; US-9587910.
(UYJO ) UNIV JOHNS HOPKIN
Lee S, MCPHETION AC;
WPI; 98-437444/37.
N-PSDB; V45823.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      active fragment. The transgenic animals have increased muscle mass and for poultry reduced cholesterol contents. Method (3) is used to treat muscle wasting or neuromuscular diseases, muscular atrophy and aging, particularly muscular dystrophy, spinal cord or traumatic injuries, congestive or obstructive lung disease. AIDS and cachexia. Method (4) is used to treat cancer of muscle, connective tissue and bone, or obesity. Also (not claimed) GDF-8 can be used to maintain myoblasts intended for transplanting or to improve efficiency of fusion. Ab can be used to detect and quantify GDF-8 (particularly in muscle, for diagnosis or monitoring), also for immunotherapy and in vivo imaging.
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W69892 star
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Ovine growth differentiation factor-8.

Growth differentiation factor-8; GDF-8; sheep; transgenic animal; transforming growth factor-beta; muscle; meat; inhibitor; obesity; neuromuscular disease; muscular dystrophy; cachexia; AIDS; cancer;
                                                                                                                                                                                                                                                                                     Transgenic animals with gene for growth differentiation factor-8 disrupted - have increased muscle and reduced choesterol contenalso use of GDF-8 inhibitors for treating cancer, obesity,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ovis
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263. .266
267. .375
/label= Mat_protein
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cc inhibiting the action of GDF-8 by treating foetal or adult muscle cc or progenitor cells with a GDF-8 inhibitor; (4) isolated nucleic acid encoding a GDF-8 protein truncated by loss of the C-terminal active fragment. The transgenic animals have increased muscle mass can for poultry reduced cholesterol contents. Method (3) is used to treat muscle wasting or neuromuscular diseases, muscular atrophy and aging, particularly muscular dystrophy, spinal cord or traumatic injuries, congestive or obstructive pinal cord or traumatic injuries, congestive or obstructive pinal cord or connective tissue and bone, or obesity. Also (not claimed) GDF-8 can be used to maintain myoblasts intended for transplanting or to improve efficiency of fusion. Ab can be used to detect and quantify GDF-8 (particularly in muscle, for diagnosis or monitoring), also for immunotherapy and in vivo imaging.
                                                                                                                                                                                                                                                                                                                                                                                          W69892
                  SEQUENCE 1.0
                                                                                                                                                                                                                                                                  101
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                                                             LYFNGKEQII YGKIPGMVVD
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                                                                                                    PKRYKANYCS
                                                                                                                                                                                  GIWQSIDVKT
                                                                                                                                          PFLEVKVTDT
                                                                                                                                                                                                                                                                  SSDGSLEDDD
                                                                                                                                                                                                                                                                                                       SRLEAIKIQI
standard;
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                                                                                                  GECEFLFLQK YPHTHLVHQA NPKGSAGPCC TPTKMSPINM
                                                                                                                                                                                                                          RPVKTPTTVF
                                                                                                                                                                                                                                                                  YHVTTETVIT
                                                                                                                                                                                  VLQNWLKQPE
                                                                                                                                          PKRSRRDFGL
Protein;
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                                                             RCGCS
                                                                                                                                        DCDEHSTESR
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                                                                                                                                          CCRYPLTVDF
                                                                                                                                                                                  DENGHDLAVT
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06-AUG-1998

05-FEB-1998; U02479.

23-MAY 1997; US-862445.

05-FEB-1997; US-795071.

28-APR-1997; US-847910.

(UTJO ) UNIV JOHNS HOPKINS SI

Lee S, MCPHERTON AC;

WPI; 98-447444/37.
Example 3; Fig 5a; 115pp; English.

This is the amino acid sequence of mouse growth differentiation factor-8 (GDF-8), a novel member of the transforming growth factor factor-8 (GDF-8), a novel member of the transforming growth factor set a superfamily that appears to relate to various cell proliferative disorders, especially those involving muscle, nerve and adipose tissue. The sequence was deduced from a cDNA clone (see V42113) isolated from a skeletal muscle cDNA library. The invention provides novel mammalian and avian GDF-8 proteins (see W69883-92). A transgenic non-human animal is claimed in which
                                                                                                                                                                                        Transgenic animals with gene for growth differentiation factor-8 disrupted - have increased muscle and reduced cholesterol contenalso use of GDF-8 inhibitors for treating cancer, obesity,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            O7-DEC-1998 (first entry)
Murine growth differentiation factor-8.
Growth differentiation factor-8; GDF-8; mouse; transgenic animal; transforming growth factor-beta; muscle; meat; inhibitor; obesity; neuromuscular disease; muscular dystrophy; cachexia; AIDS; cancer; neuromuscular disease; muscular dystrophy; cachexia; AIDS; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cleavage_site
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                                                                                                                                                                                                                                                                                                                                                                                                                                        /note- "Asn is N-glycosylated"
264. 267
268. 376
/label- Mat_protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers 72. .74
                                                                                                                                                                                                                                                                                                           SCHOOL MEDICINE
                                                                                                                                                                                                           contents
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CC are: (1) chicken or turky eggs or meat, beef, milk, pork and lamb CC from these animals; (2) method for increasing muscle mass in animals by administering an antibody (Ab) that binds to GDF-8; (3) cc inhibiting the action of GDF-8 by treating foetal or adult muscle cor progenitor cells with a GDF-8 inhibitor; (4) isolated nucleic active fragment. The transgenic animals have increased muscle mass in condition of GDF-8 protein truncated by loss of the C-terminal correct fragment. The transgenic animals have increased muscle mass correct muscle wasting or neuromuscular diseases, muscular atrophy and aging, particularly muscular dystrophy, spinal cord or traumatic injuries, congestive or obstructive ing disease, AIDS cand cachexia, Method (4) is used to treat cancer of muscle, and connective tissue and bone, or obesity. Also (not claimed) GDF-8 can be used to maintain myoblasts intended for transplanting or to improve efficiency of fusion. Ab can be used to detect and calse of immunotherapy and in vivo imaging.
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06-AUG-1998: U02479.
23-MAY-1999: U0-862445.
23-MAY-1997: US-862445.
05-FEB-1997: US-847910.
28-APR-1997: US-847910.
(UTJO) UNIV JOHNS HOPKINS SI
Lee S, McPherron AC;
WPI; 98-437444/37.
N-PSDB: V45813.
N-PSDB: V45813.
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07-DEC-1998 (first entry)
Human growth differentiation factor-8.
Human growth differentiation factor-8; human; transgenic acrowth differentiation factor-8; muscle; meat; inhibitor; or neuromuscular disease; muscular dystrophy; cachexia; AIDS;
Transgenic animals with gene for growth differentiation factor-8 disrupted - have increased muscle and reduced cholesterol contents, also use of GDF-8 inhibitors for treating cancer, obesity, neuromuscular disease Example 3; Fig 5c: 125pp; English.

This is the amino acid sequence of human growth differentiation factor-8 (GDF-8), a novel member of the transforming growth factor-beta superfamily that appears to relate to various cell proliferative disorders, especially those involving muscle, nerve
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Protein
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Homo sapiens.
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263. .266
267. .375
/label= Ma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   animal;
obesity;
cancer;
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8888888888888888888888888888888888
CC and adipose tissue. The sequence was deduced from a CDNA clone CC (see v4580) isolated from a skeletal muscle CDNA library. The CC invention provides movel mammalian and avian GDP-8 proteins (see GC are: (1) chicken or turkey eggs or meat, beef, milk, pork and lamb CC are: (1) chicken or turkey eggs or meat, beef, milk, pork and lamb CC animals by administering an antibody (Ab) that binds to GDP-8; (3) CC inhibiting the action of GDP-8 by treating foetal or adult muscle cor progenitor cells with a GDP-8 by treating foetal or adult muscle caid encoding a GDP-8 protein truncated by loss of the C-terminal cc acid encoding a GDP-8 protein truncated by loss of the C-terminal cc and for poultry reduced cholesterol contents. Method (3) is used to treat muscle wasting or neuromuscular diseases, muscular artrophy CC and adjng, particularly muscular dystrophy, spinal cord or traumatic injuries, congestive or obstructive lung disease, AIDS cand cachexia. Method (4) is used to treat cancer of muscle, and cachexia. Method (4) is used to treat cancer of muscle. Cc can eachexia with a GDP-8 protein truncated by a consective tissue and bone, or obstructive lung disease, AIDS (and cachexia, Method (4) is used to treat cancer of muscle. Cc can be used to maintain myoblasts intended for transplanting or to CC improve efficiency of fusion. Ab can be used to detect and cc calso for immunotherapy and in vivo imaging.
                                                                                   monitoring),
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W69885 Length: 375 February 11, 2000 15:49 Type: P 101 351 301 201 51 -LYFNGKEQII PKRYKANYCS PFLEVKVTDT GIWQSIDVKT VVKAQLWIYL SSDGSLEDDD SRIEAIKIQI LSKLRLETAP NISKDVIRQL LPKAPPLREL IDQYDVQRDD MOKLQLCVYI YLFMLIVAGP VDLNENSEQK ENVEKEGLON YGKIPAMVVD GECEFVFLQK PKRSRRDFGL VLQNWLKQPE RPVETPTTVF YHATTETIIT YPHTHLVHQA DCDEHSTESR RCGCS SNLGIEIKAL VQILRLIKPM MPTESDFLMQ NPRGSAGPCC CCRYPLIVDE DENGHDLAVT KDGTRYTGIR VDGKPKCCFF TPTKMSPINM EAFGWDWIIA SLKLDMNPGT KESSKIQYNK ACTWRQNTKS Check: 1814

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WO9833887-A1.

PN W09833887-A1.

PN W09833887-A1.

PN 05-EBB-1998; U02479.

PR 05-EBB-1998; US-799071.

PR 28-APR-1997; US-847910.

PR 28-APR-1997; US-847910.

PR 28-APR-1997; US-847910.

PR 198-1997; US-847910.

PR 198-1997; US-847910.

PT 128-25, McPherron Ac.;

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W69886 stand
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Protein
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Babboon growth differentiation factor-8.

Growth differentiation factor-8; GDF-8; babboon; transgenic growth differentiation factor-beta; muscle; meat; inhibitor; o transforming growth factor-beta; muscle; meat; albis; neuromuscular disease; muscular dystrophy; cachexia; AIDS;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    therapy.
Papio sp.
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263. .266
267. .375
/label= Mat_protein
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  obesity;
; cancer;
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CC proliferative disorders, especially those involving muscle, nerve
CC and adipose tissue. The sequence was deduced from a cDNA clone
CC (see V45817) isolated from a skeletal muscle cDNA library. The
CC invention provides novel mammalian and avian GDF-8 proteins (see
CC GDF-8 expression is disrupted or incerfered with. Also claimed
CC are: (1) chicken or turkey eggs or meat, beef, milk, pork and lamb
CC arimals by administering an antibody (Ab) that binds to GDF-8; (3)
CC inhibiting the action of GDF-8 by treating foretal or adult muscle
CC or progenitor cells with a GDF-8 by treating foretal or adult muscle
CC acid encoding a GDF-8 protein truncated by loss of the C-terminal
CC acid encoding a GDF-8 protein truncated by loss of the C-terminal
CC and for poultry reduced cholesterol contents. Method (3) is used
CC to treat muscle wasting or neuromuscular disease, muscular atrophy
CC and aging particularly muscular dystrophy, spinal cord or
CC tranmatic injuries, congestive or obstructive lung disease, AIDS
CC and cachexia. Method (4) is used to treat cancer of muscle,
CC and cachexia. Method (6) is used to treat cancer of muscle,
CC and cachexia method (7) is used to treat cancer of muscle,
CC and cachexia. Method (7) is used to treat cancer of muscle,
CC and cachexia method (7) is used to treat cancer of muscle,
CC and cachexia of the muscle, for diagnosis or monitoring),
CC also for immunotherapy and in vivo imaging.
CC also for immunotherapy and in vivo imaging.
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06-AUG-1998.
05-FEB-1998; U02479.
23-MAY-1997; US-862445.
05-FEB-1997; US-95071.
28-APR-1997; US-847910.
(UYJO) UNIV JOHNS HOPKINS S.
Lee S. MCPherion AC;
WPI, 98-437444/37.
Transgenic animals with gene for growth differentiation factor-8 disrupted - have increased muscle and reduced cholesterol contents, also use of GDF-8 inhibitors for treating cancer, obesity, neuromuscular disease Example 9; Fig 14b; 125pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              07-DEC-1998 (first entry)
Bovine growth differentiation factor-8,
Growth differentiation factor-8; GDF-8; human; transgenic animal;
transforming growth factor-beta; muscle; meat; inhibitor; obesity;
neuromuscular disease; muscular dystrophy; cachexia; AIDS; cancer;
therapy.
                                                                                                                                                              WPI; 98-437444/37.
N-PSDB; V45818
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07-DEC-1998
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263. .266
267. .375
/label= Mat_protein
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CC factor-8 (GDF-8), a novel member of the transforming growth differentiation CC factor-8 (GDF-8), a novel member of the transforming growth factor-CC beta superfamily that appears to relate to various cell CP proliferative disorders, especially those involving muscle, nerve CR and adjose tissue. The sequence was deduced from a cDNA clone CR (see v45818) isolated from a skeletal muscle cDNA library. The CR invention provides novel mammalian and avian GDF-8 proteins (see CR 65883-92). A transgenic non-human animal is claimed in which CR GDF-8 expression is disrupted or interfered with. Also claimed CR creit (1) chicken or turkey eggs or meat, beef, milk, pork and lamb CR are: (1) chicken or turkey eggs or meat, beef, milk, pork and lamb CR are these animals: (2) method for increasing muscle mass in CR animals by administering an antibody (Ab) that binds to GDF-8; (3) inhibiting the action of GDF-8 by treating foetal or adult muscle call encoding a GDF-8 protein truncated by loss of the C-terminal CR and for poultry reduced cholesterol contents. Method (3) is used to treat muscle wasting or neuromuscular diseases, muscular atrophy CR and aging, particularly muscular dystrophy, spinal cord or traumatic hijuries, congestive or obstructive lung disease, AIDS CR and cachexia. Method (4) is used to treat cancer of muscle. CR can be used to maintain myoblasts intended for transplanting or to cambertive tissue and bone, or obesity. Also (not claimed) GDF-8 CR improve efficiency of fusion. Ab can be used to detect and cachect and duantify GDF-8 (PATKicularly in muscle, for diagnosis or monitoring), sequence 375 AA; Sequence 375 ₽Ą,

W69887 Length: 375 February 11, 2000 15:49 Type: P Check: 9305

51 SRLEAIKIQI MORLQISVYI LSKLRLETAP NISKDAIRQL YLFMLIVAGP VDLNENSEQK ENVEKEGLCN LPKAPPLLEL IDQFDVQRDA ACLWRENTTS

101 SSDGSLEDDD YHARTETVIT MPTESDLLTQ VEGKPKCCFF

151 LVKAQLWIYL RPVKTPATVE VQILRLIKPM KDGTRYTGIR SLKLDMNPGT

201

VLQNWLKQPE

SNLGIEIKAL

DENGHDLAVT

251 PFLEVKVTDT PKRSRRDFGL DCDEHSTESR CCRYPLIVDF EAFGWDWIIA

PKRYKANYCS GECEFVFLQK YPHTHLVHQA NPRGSAGPCC

351 301

LYFNGEGQII YGKIPAMVVD

RCGCS

_SEQUENCE 1.0 W80799 stand standard; Protein; 244 A

710248578-A.
22-581-1998; 065618.
05-MAR-1997; JP-065618.
(NITT) NITTO CHEM IND C
WPI: 98-560730/48. W80799;
08-DEC-1998 (first entry)
08-DEC-1998 (first entry)
Rhodococcus nitrile hydratase gene fragment product.
Rhodococcus; nitrile hydratase gene; expression vector; control factor; nitrilase gene promoter.
Rhodococcus erythropolis.

V65530 ၀

Expression vector - for Rhodococcus genus microbe claim 2; Pages 5-6; 10pp; Japanese.

This represents a product of the nitrile hydratase gene fragment used in the construction of an expression vector for a Rhodococcus genus microbe. The expression vector contains a DNA region encoding a control factor having the ability of activating a nitrilase gene promoter; a DNA region of the nitrilase gene promotor activated by the above control factor; a DNA region producible in the cell of a Rhodococcus genus microbe, and a chemically resistant DNA region functioning in the Rhodococcus genus microbe. An exotic gene can be expressed constitutionally.

W80799

Length:

244

February

11,

2000 15:49

Type:

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Check: 2018

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ID W80624 standa
                                                                                                                                                                                                                                                                                                                                                                             This sequence represents a Streptococcus pneumoniae GTP-binding CC protein. The invention provides DNA sequences (V65201 to V65304) from CC protein. The invention provides DNA sequences you follow the Streptococcus pneumoniae genome and corresponding protein sequences (W80605 to W80728). The protein sequences are classified as hypothetical, CC cell wall biosynthetic, external target, or minimal gene set proteins. A CC recombinant host containing a vector comprising any of the above nucleic carids can be used for the recombinant expression of the proteins. The CC acids can be used for the recombinant expression of the proteins. The CC invention also provides a DNA chip having arrayed on it at least 15 base CC invention also provides a DNA chip having arrayed on it at least 15 base CC inventions for evaluating gene expression in S. pneumoniae and for CC identifying virulence genes in S. pneumoniae. Antibodies that selectively bind to the above proteins or peptide fragments can be used to treat CC S. pneumoniae infection. The antibodies can also be used to detect S. pneumoniae cells.
ID AC DT DE KW KW KW PD PD PD PD
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(ELIL) LILLY & CO ELI.
Baltz RH. Burgett SG, Dehoff BS, Hoskins JA, Jaskunas Baltz RH. Burgett SG, Dehoff BS, Rockey PK, Rosteck PR, Mills BJ, Norris FH, Peery RB, Rockey PK, Rosteck PR, Mills BJ, Norris FH, Peery RB, Rockey PJ, Treadway PJ, Skatrud PL, Smith MC, Solenberg PJ, Treadway PJ,
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09-DEC-1997; U22578.
13-DEC-1996; US-036281.
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WPI; 98-348529/30.
N-PSDB; V65202.
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                                                                                                                                                           SEQUENCE 1.
W68494 star
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                                                                 W68494;
07-JAN-1999 (first entry)
E2 papillomavirus protein;
E2 protein; apoptosis; HPV
cervix; virus-infected cel
                                                                                                                                                                                                                                            101
                                Papillomavirus.
WO9832861-A1.
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                                                                                                                                                             Protein;
                                                                     ein encoded by plasmid pCGE2.18VoNCO.
HPV; infection; papillomavirus-associated
cell; p53 tumour repressor.
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                                                                                                 cancer;
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29-JAN-1998; F00169

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RAGE REGERENCE OCCOONS RAGER REGERENCE REGEREN
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PP 11-NOV-1998.

PP 11-NOV-1998.

303597.

PF 01-MAY-1998; US-071342.

PR 01-MAY-1999; US-071342.

PR 02-MAY-1999; US-045890.

PR 02-MAY-1999; US-04589.

PR 02-MAY-1998; US-04589.

PR 02-MAY-1998; US-04589.

PR 02-MAY-1998; US-04-MAY-19689.

PR 02-MAY-1999; US-04-MAY-19689.

PR 02-MAY-1998; US-04-MAY-19689.

PR 02-MAY-1999; US-04-MAY-19689.

PR 02-MAY-1998; US-04-MAY-19689.

PR 02-MAY-1998; US-04-MAY-19689.

PR 02-MAY-1998; US-04-MAY-19689.

PR 02-MAY-1996; US-04-MAY-19689.

PR 02-MAY-19689.

PR 02-MAY-1996.

PR 02-MAY-1996.

PR 02-MAY-19689.

PR 02
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Disclosure; Fig 12; llipp; French.

Disclosure; Fig 12; llipp; French.

The present sequence is encoded by an insert of plasmid pCGE2.18VoNCO represent sequence is encoded by an integrated part of the human to induce apoptosis in cells that have integrated part of the human to induce apoptosis in cells that have integrated part of the human to induce apoptosis in cells that have integrated part of the human to including those expressing wild-type E2) and the corresponding vectors (including those expressing wild-type E2) and the corresponding vectors or truncated E2 (E2TR) are all useful for treating or preventing proteins or truncated E2 (E2TR) are all useful for treating or preventing papillomavirus infection, particularly papillomavirus-associated cancers papillomavirus december of the cervix uteri). The proteins, and sequences expressing them, also induce apoptosis of virus-infected cells and increasing the activity of the p53 tumour repressor.

Sequence 368 AA;
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(INSP ) INST PASTEUR.
(UYME-) UNIV MEXICO NACIONAL AUTONOMA.
Demotret C, Desaintes C, Goyat S, Thierry
WPI: 98-427957/36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EPRGI: EPO primary response gene 1; diagnosis; gene therapy; immunity; disease; vaccine; inoculate; antibody; T cell; anaemia; polycythemia; cancer; neutropenia; AIDS; diabetes, myelosuppression; allergy; asthma; autoimmune disease; inflammatory disease; chromosome mapping; human. Homo sapiens.

EP-877030-A2.
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protein #1.
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PT New EPO primary response gene polypeptides and polynucleotides - PT useful as diagnostic reagents and for prevention and treatment of PT useful as diagnostic reagents and for prevention and treatment of PT cancer and autoimmune and inflammatory diseases
PS Claim ; Page 2; 25pp; English.

CC (EPRG1) polypeptide. EPRG1 polypeptides and polynucleotides are useful CC (EPRG1) polypeptide. EPRG1 polypeptides and polynucleotides are useful CC (EPRG1) polypeptides on the EPRG1 not polypeptide or mRNA expression levels can the used to screen for agonists and antagonists CC EPRG1 polypeptide by measuring resulting mRNA levels with CC ELISA. These can be used in treatment to activate (agonist) or inhibit CC (antagonist eg EPRG1 polypeptide by measuring resulting mRNA levels with CC (antagonist eg EPRG1 ligand, receptor or substrate) EPRG1 activity, in CC addition to direct administration of antisense sequences to prevent CC expression, or EPRG1 polypeptides to treat conditions associated with CC a lack of EPRG1 protein. Gene therapy may also be used to affect CC endogenous EPRG1 polypeptide production. EPRG1 antibodies are useful for collecting an immune response to immunise and prevent diseases, and for conduction to inoculate against diseases by inducing an antibody and T-cell CC expression, autoimmune diseases, rheumatoid artivitis and multiple CC expression, autoimmune diseases, including asthma and allergies. The EPRG1 polypeptide is also useful for mapping the gene to a chromosome,
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EPRG1; EPO primary response gene 1; diagnosis; gene therapy; immunity; disease; vaccine; inoculate; antibody; T cell; anaemia; polycythemia; cancer; neutropenia; AIDS; diabetes, myelosuppression; allergy; asthma autoimmune disease; inflammatory disease; chromosome mapping; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Dillon S, Lord K;
WPI; 98-570499/49.
N-PSDB; V69308.
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07-MAY-1998; 303597.
01-MAY-1998; US-071342.
07-MAY-1997; US-045890.
(SMIK ) SMITHKLINE BECKMAN CORP.
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W81740

Length:

120

February 11,

2000 15:49

Type: ъ

Check: 3575

SEQUENCE 1.0 W72902 standard; Protein; 226

AA.

21-JAN-1999 (first entry)

Mycobacterium tuberculosis; antigen CFP19A.
Mycobacterium tuberculosis; antigen; vaccine; immunological;
immunogen; infection.

101

51

QKLEEENPEA

AQALRAKIES ENPDAARADR CVSPTSQARD ARRPLARSAR

AVQTEDKYGV KIPDEDLAGL RTVGDVVAYI

TPEKSFVDDL

DIDSLSMVEI

TOS SERVICE TO SERVICE

Peptide

1. .38 /label- signal Location/Qualifiers

Mycobacterium tuberculosis.

W09844119-A1. 08-OCT-1998.

01-APR-1998; DK0132. 05-JAN-1998; US-070448. 02-APR-1997; DK-000376. 18-APR-1997; US-044624. 10-NOV-1997; DK-001277. (STAT-) STATEMS SERUM INST. Andersen P, Florio W, Nielsen R

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Oettinger

PB

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                                                 PR 13-MAR-1997; U18293.

PR 11-MAR-1997; US-938112.

PR 11-CT-1996; US-730510.

PR (CORI-) CORIXA CORP.

PR (CORI-) CORIXA CORP.

PI Reed SG, Skeiky YAW, Twardzik DR, Vedvick TS;

PI Reed SG, Skeiky YAW, Twardzik DR, Vedvick TS;

PI Reed SG, Skeiky YAW, Twardzik DR, Vedvick TS;

PI Reed SG, Skeiky YAW, Twardzik DR, Vedvick TS;

PI Reed SG, Skeiky YAW, Twardzik DR, Vedvick TS;

PR N-PSDB; V64551.

PR N-P
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ID W81740 standard; Protein; 120
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M. tuberculosis immunogenic polypeptide RDIF5.

Tuberculosis; immunogenic; soluble; antigen; protective immunity; vaccine; pharmaceutical; infection; diagnosis.
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Sequence
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                                   diagnosis of
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              120
                                          tuberculosis.
              8
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ul to screen for
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STTSALTPGP AAAAADAAAS

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W72902
                              Py Vascular endothelial growth factor-B iso:forms, and DNA encoding provided the provided the provided to inducing anglogenesis and cellular proliferation, provided the provided to inducing anglogenesis and cellular proliferation, provided the present sequence represents human vascular endothelial growth factor the present sequence represents human vascular endothelial growth factor the provided the proliferation of conditional cells and for stimulating anglogenesis (the proliferation of new capillaries form pre-existing blood vessels). These activities are useful for treating tissue growth and repair, including organic repair. This is also useful in pregnancy, in follicle development, as these processes must occur in development of the placenta. The proteins can also be used to raise antibodies, either for use in the proteins of the proteins or as inhibitors of their action. This is capacially useful as angiogenesis is required by tumours as they need consumed to the proteins or as inhibitors of their action. This is sepacially useful as angiogenesis is required by tumours as they need consumed to the proteins or as inhibitors of the proteins or as a subject to grow and proliferate.
W80495
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24 NOV-1998.
01-MAR-1996; US-609443.
01-MAR-1996; US-597651.
06-DUC-1995; US-469427.
06-DEC-1995; US-59063.
(LUDW-) LUDWIG INST CANCER RES.
(LUDW-) UNIV HELSINKI LICENSING LTD OY
Alitalo K, Eriksson U, Olofsson B, Pa)
NUTSEEP V65560
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WPI; 98-542705/
N-PSDB; V63926
New isolated my
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Human vascular endothelial
Vascular endothelial growt
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207
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ial growth factor; VEGF; proliferation; human;
angiogenesis; tissue growth; organ repair.
  February 11,
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ID W85306 standard;
AC W85306;
www.i. yo.44/0/y.jo.

Composition containing peptide - can bind to human leucocyte antigen
promposition containing peptide - can bind to human leucocyte antigen
promposition containing peptide - can bind to human leucocyte antigen
promposition containing peptide - can bind to human leucocyte antigen
promposition containing peptides, which can bind to
CC w85284-451 represent helper T-cell class II peptides, which can bind to
CC the human leucocyte antigens (HLA) DRAW4, DRI and DR7. The peptides
CC are used in the course of the invention. The specification describes
CC are used in the course of the invention. The specification describes
CC the treatment or prevention of viral, fungal, bacterial or parasitic
CC infections (e.g. hepatitis, acquired immune deficiency syndrome or
CC malaria) or cancer (e.g. renal or cervical carcinoma, lymphoma, prostate
CC cancer or condyloma acuminatum). Helper T-cell peptides may be used
CC alone to induce a helper T-cell response, e.g. in cases of autoimmune
CC disease, allograft rejection, allergy, Lyme disease, hepatitis,
CC hypersensitivity.
SQ Sequence 24 AA;
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30-JUL-1998; U01373.
23-JAN-1998; U01373.
07-FEB-1997; US-037432.
23-JAN-1997; US-036713.
(EPIM-) EPIMMUNE INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Helper T-cell class II peptide derived from mouse invariant chain. Helper T-cell peptide; human leucocyte antiqen; HLA; DR444; DR1; Helper T-cell peptide; human leucocyte antiqen; HLA; CR444; DR1; DR7; cytotoxic T lymphocyte; CTL; hepatitis; autoimmune disease; acquired immune deficiency syndrome; malaria; cancer; allograft rejection; allery; Lyme disease; hepatitis; post-streptococcal endocarditis; glomerulonephritis;
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WPI; 98-427679/36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   southwood S;
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W85306
               Length: 24
LPKPPKPVSK MRMATPLLMQ ALPM
                 February 11, 2000
                   15:49
                   Type:
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                    Check:
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W86202;
W86202;
Hindan VEGF-related factor (VRF)-2 sequence.
Hindan VEGF-related factor (VRF)-2 sequence.

**WEGF: VRP; vascular endothelial growth factor; VEGF-related protein; recombinant; truncated; gene therapy; anglogenesis; cardiac ischaemi recombinant; collateral vessel development; cell growth; migration; heatorearly collateral vessel development; vascular disease; intesting lower limb ischaemia; stroke; peripheral vascular disease; intesting wound healing; skin; VEGF-related factor; VRF; vascular permeability
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               _SEQUENCE 1.0
W86202 standard;
(COLL-) COLLATERAL THERAPBUTICS.
Bohlen 9:
WPI; 99-009426/01.
WPI; 99-009426/01.
New truncated vascular endothelial growth factor-related protein subunits - lack part of the N-terminal sequence, used to stimulate angiogenesis, e.g. for treating heart disease and ischaemia pisclosure; Fig 1; 113pp; English.
Disclosure; Fig 1; 113pp; English.
This represent the amino acid sequence of human VRF-2 (a vascular represent the amino acid sequence factor). The invention
                                                                                                                                                                               20-APR-1998; U07801.
25-APR-1997; US-842984.
(COLL-) COLLATERAL THEI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     protein;
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101

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MGTVAKQLVP AALLQLAPAQ

SCYTYQRCGG APVSQPDAPG

CCPDDGLECV PTGQHQVRMQ

HORKVVSWID

VYTRATCOPR

GVGSLSQSTI

CRFESLTLSH NNMIALKPIL

QAWLEEAEGA QREKMNKPEI

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                                                                                                                                                                                                                                                                                                                                                                                      New diagnosis of leukaemia - by detecting genes for tumour antigen rejection precursors or corresponding proteins
Claim 11, Pages 88-69, 88pp; English.
The present sequence represents the amino acid sequence of the SIAX DP2-64 (Oct-T1) gene. The protein is a tumour rejection antigen precursor (TRAP). The specification describes the treatment of disorders which characterised by expression of a leukaemia-associated nucleic acid such as TRH. The products are used for in vivo or in vitro screening for leukaemia, lymphoma or other cancers by usual hybridisation/amplification or immunoassay methods. TRAPs, when molecules, or nucleic acid encoding them, are useful in vaccines for the processed to antigens or complexed with HIA (human lymphocyte antigen) molecules, or nucleic acid encoding them, are useful in vaccines for
                                                                                                      151
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22-APR-1998; U07784.
25-APR-1997; US-845998.
(LUDW-) LUDWIG INST CANCER RES.
Boon-Falleur T, Coulie PG, De Smet C, Lucas S, Van Baren WPI; 99-09425/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12-FEB-1999 (first entry)
Amino acid sequence of the SIAX DP2-64 (Oct-T1) gene.
Tumour rejection antigen precursor; TRAP; TRH; leukaemia;
Lymphoma; cancer; HLA; human lymphocyte antigen; vaccine;
SIAX DP2-64; Oct-T1.
Homo sapiens
Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                          treating leukaemia.
Sequence 420 AA;
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W68541 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             provides truncated VRP (VEGF-related protein) subunits that have at least one amino acid N-terminal to the first Cys of the core sequence deleted. Host cells transformed or transfected with expression vectors containing nucleic acids encoding the truncated VRP subunits are used to produce the truncated proteins recombinantly. The truncated VRP subunits, and in vitro anglogenic activity and are used to stimulate angiogenesis particularly coronary collateral vessel development in cases of cardiac ischaemia; to disease; to treat ischaemia (e.g. cardiac, chronic coronary or chronic lower limb ischaemia; stroke and peripheral vascular disease); to promote healing of wounds (of skin or intestines), and to increase vascular coronary coular coronary coronar
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VGAAGLASIC DSDTDPRELE
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                                                                                          GPGGGGGGPG
                                                                                                                                               SSPSLALMAG
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IKLGVTQADV GSALANLKIF

Homo sapiens. WO9849300-A2.

05-NOV-1998.
20-APR-1998; U07801.
25-APR-1997; US-842984.
(COLL-) COLLATERAL THERAPEUTICS
Bohlen P;

16-FEB-1999 (first entry)
Human VRF-2 truncated fragment 1.
VEGF, VRP, vascular endothelial growth factor; VEGF-related protein; recombinant; truncated; gene therapy; anglogenesis; cardiac ischaemia; coronary; collateral vessel development; cell growth; migration; heart lower limb ischaemia; stroke; peripheral vascular disease; intestine; wound healing; skin; VEGF-related factor; VRF; vascular permeability.

SEQUENCE 1.0

86214 standard; protein;

AA.

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PT New truncated vascular endothelial growth factor-related protein PT subunits - lack part of the N-terminal sequence, used to stimulate PT subunits - lack part of the N-terminal sequence, used to stimulate PT angiogenesis, e.g. for treating heart disease and ischaemia Claim 5; Fig 2B; 113pp, English.

CC The invention relates to truncated VRP (vascular endothelial growth CC factor (VEGF)-related protein) subunits that have at least one amino acid CC reminal to the first Cys of the core sequence deleted. Host cells CC truncated or transfected with expression vectors containing nucleic CC truncated proteins recombinantly. The truncated VRP subunits are used to produce the CC expressed from gene therapy vectors, have in vivo and in vitro angiogenic CC collateral vessel development in cases of cardiac ischaemia; to stimulate angiogenesis, particularly coronary CC endothelial cell growth and migration in vitro; to treat heart disease; CC terest ischaemia (e.g. cardiac, chronic coronary or chronic lower limb vounds (of skin or intesthnes), and to increase vascular permeability of Sequences W86234 to W86239 represent full length VRP sequences from Sequence 185 AA;
                                                                                                                                                             W86235
       151
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16-FEB-1999 (first entry)
16-FEB-1999 (first entry)
Human VRF (VEGF-related factor)-2 full length sequence.
VEGF; VRF; vascular endothelial growth factor; VEGF-related protein; recombinant; truncated; gene therapy; angiogenesis; cardiac ischaemia; coronary; collateral vessel development; cell growth; migration; heart; lower limb ischaemia; stroke; peripheral vascular disease; intestine; wound healing; skin; vascular permeability; VRF.
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05-APR-1998: U07801.
25-APR-1997: US-842984.
25-APR-1997: US-842984.
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WO9849300-A2.
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GPSAHAAPST TSALTPGPAA
                                                                                                                                                        Length: 185
                                     CECRPKKDSA VKPDRAATPH
                                                                             CVTVQRCGGC
                                                                                                                     PVSQPDAPGH
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                                                                             CPDDGLECVP
                                                                                                                QRKVVSWIDV
                                                                                                                                                 February 11, 2000 15:49 Type: P Check: 8227
  AAADAAASSV
                                     HRPQPRSVPG
                                                                      TGQHQVRMQI LMIRYPSSQL GEMSLEEHSQ
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                              WDSAPGAPSP ADITHPTPAP
                                                                                                              VVVPLTVELM
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!!AA_SEQUENCE 1

standard;

protein; 168

151 101

ALTPGPAAAA ADAAASSVAK

PDRAATPHHR PQPRSVPGWD SAPGAPSPAD

ITHPTPAPGP SAHAAPSTTS

DDGLECVPTG QHQVRMQILM IRYPSSQLGE MSLEEHSQCE CRPKKDSAVK

51

16-FEB-1999 (first entry)
16-FEB-1999 (first entry)
Human VRF-2 truncated fragment 3.
VEGF- VRPy vascular endothelial growth factor; VEGF-related protein;
recombinant; truncated; gene therapy; angiogenesis; cardiac ischaemia.
coronary; collateral vessel development; cell growth; migration; hear
lower limb ischaemia; stroke; peripheral vascular disease; intestine;
wound healing; skin; VEGF-related factor; VRF; vascular permeability.

protein; ischaemia;

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CC That is Fig 2B; 113pp; English.

CC The invention relates to truncated VRP (vascular endothelial growth CC factor (VEGF) related protein) subunits that have at least one amino acid CC transformed or transfected with expression vectors containing nucleic CC cacids encoding the truncated VRP subunits are used to protein cc the cc expressed from gene therapy vectors, have in vivo and in vitro anglogenic CC collateral vessel development in cases of cardiac ischaemia; to stimulate endoughers to treat ischaemia (e.g. cardiac, chronic coronary or chronic lower limb CC ischaemia; stroke and peripheral vascular disease); to promote healing of CC wounds (of skin or intestines), and to increase vascular permeability.

CC sequences W86214 to W86217 represent truncated fragments of VRF-2 (VEGF-CC related factor).
                                                                   PI Bohlen P;
Boh
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Homo sapiens.
WO9849300-A2.
05-NOV-1998.
20-APR-1998; U07801.
25-APR-1997; US-842984.
25-APR-1997; US-842984.
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W86215 st
W86215;
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Human VRF-2 truncated fragment 2.
VEGF. VRP, vascular endothelial growth factor; VEGF-related protein; recombinant; truncated; gene therapy; anglogenesis; cardiac ischaemia; coronary; collateral vessel development; cell growth; migration; heart lower limb ischaemia; stroke; peripheral vescular disease; intestine; wound healing; skin; VEGF-related factor; VRF; vascular permeability.
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e 178 AA;
    factor).
e 173 AA;
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05-NOV-1998. 20-APR-1998; U07801. 25-APR-1997; US-842984. (COLL-) COLLATERAL THERAPEUTICS.

PI Bohlen P;
Boh

W86216 Length: 168 February 11,

IDVYTRATCQ

PREVVVPLTV

ELMGTVAKQL 2000 15:49

VPSCVTVQRC GGCCPDDGLE Type: P

Check: 2203

Sequence

168 AA;

101

51

CVPTGQHQVR MQILMIRYPS SQLGEMSLEE HSQCECRPKK DSAVKPDRAA

VPGWDSAPGA PSPADITHPT PAPGPSAHAA PSTTSALTPG

151

PAAAAADAAA SSVAKGGA

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AA!
       PRESENTATION OF THE PROPERTY O
05-NOV-1998.

20-APR-1998; U07801

25-APR-1997; US-8429

(COLL-) COLLATERAL 'Bohlen P;
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W86217 standa
                                                                                                                                                                                                                                                                                                                                                                                                                                        coronary; collateral v
lower limb ischaemia;
wound healing; skin; v
                                                                                                                                                                                                                                                                                                                                                                                                                                 16-FEB-1999 (first entry)
Human VRF-2 truncated fragment 4.
VEGF; VRP; vascular endothelial growth factor; VEGF-related protein; recombinant; truncated; gene therapy; anglogenests; cardiac ischaemia; coronary; collateral vessel development; cell growth; migration; heart lower limb ischaemia; stroke; peripheral vascular disease; intestine; wound healing; skin; VEGF-related factor; VRF; vascular permeability.
                                                                                                                                                                                                                                                                                                                            Homo sapiens.
W09849300-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             standard; protein; 163
                                                                                    THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   migration; heart;
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W86215

Length: 173

February 11,

2000 15:49

Type: P

Check: 4363

KVVSWIDVYT RATCOPREVV VPLTVELMGT VAKQLVPSCV TVQRCGGCCP

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IIAA_SEQUENCE 1.0

IN W85119 stand

AC W85119;

DT 11-FEB-1999

DE A delta-5 ele

KW Fatty acid;

KW cancer; dishe

KW cholesterol

KW cholesterol

KW cholesterol

KW cholesterol

KW myalgic ence

KW acute respin

OS Mortierella

PN W09846765-Al

PD 22-CCT-1998;

PF 11-APR-1998;

PF 11-APR-1998;

PR 11-APR-1998
DR N-PSDB; V83028.

PT New nucleic acid encoding delta5 and other desaturase enzymes - PT useful in production of oils of increased arachidonic acid content, produced, e.g. for treating cancer, as foods, animal feeds and cosmetics Claim 3; Fig 3A-D; 153pp; English.

CC The present sequence represents a Montierella alpina fatty acid delta-5 cd desaturase enzyme. The enzyme sequence is used in the methods of the invention. The specification describes methods for desaturating a cfatty acid and for producing a desaturated fatty acid by expressing compositions and included the enzyme can be used for desaturating fatty acid molecule. The enzyme can be used for desaturated fatty acid so the introduces a double bond carbons 5 and 6 from the carboxyl end of a fatty acid molecule. The enzyme can be used for desaturated fatty acids. The enzyme can be used to produce polyunsaturated fatty acids. The carbox, in cosmetics or in animal feed. The polyunsaturated fatty acids can be used for treating malnutrition, in pharmaceutical compositions, in cosmetics or in animal feed. The polyunsaturated fatty acids can be used for treating e.g. restenosis after angioplasty, concer, diabetes or recame or reduce blood pressure. They can also be used to inhibit compositions are approached any produce polyunsaturated fatty acids or ezema or reduce blood pressure. They can also be used to inhibit compositions of vessel wall smooth muscle and fibrous tissue,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PR WPI; 99-009426/01.

PT New truncated vascular endothelial growth factor-related protein PT subunits - lack part of the N-terminal sequence, used to stimulate PT subunits - lack part of the N-terminal sequence, used to stimulate PT angiogenesis, e.g. for treating heart disease and ischaemia SC laim 5; Fig 2B; 113pp; English.

CC The invention relates to truncated VRP (vascular endothelial growth CC Invention relates to truncated VRP (vascular endothelial growth CC Invention relates to truncated VRP subunits that have at least one amino acid CC N-terminal to the first Cys of the core sequence deleted. Host cells CC cacids encoding the truncated VRP subunits are used to produce the CC truncated proteins recombinantly. The truncated VRP subunits, optionally CC expressed from gene therapy vectors, have in vivo and in vitro angiogenic CC collateral vessel development in cases of cardiac ischaemia; to stimulate angiogenesis, particularly coronary CC collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac ischaemia; to stimulate collateral vessel development in cases of cardiac 
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22-OCT-1998. U07422.
11-APR-1997. US-833610.
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) ABBOTT LAB.
( ) CALGENE LLC.
( ) S. Huang Y, Knutzon
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                                                                                                                                                                                                                                                                                                                                            reduce or prevent gastro-intestinal bleeding and other side effects caused by non-steroidal anti-inflammatory drugs, prevent or treat endometriosis and premenstrual syndrome, treat myalgic encephalomyelitis and chronic fatigue after viral infections, treat AIDS, multiple sclerosis, acute respiratory syndrome, hypertension and inflammatory skin disorders.
                              SEQUENCE 1
W73507 star
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                                                                   YPDILAIIKN TCSEYKVPYL
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                                                                                                                                                                                                                                                                                MGTDQGKTFT WEELAAHNTK
                           standard; Protein;
                                                                                                                                                                                                                                                                                                                                    446 AA;
(first entry)
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                                                                                                                                                                                                  ACAQVGLNPL
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                                                                 VKDTFWQAFA
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                                                                 SHLEHLRVLG
                                                                                           TGSLNYQAVH
                                                                                                                    LALTFQANHV
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ID W73507 standard; Protein; 317 AA.

AC W73507; standard; Protein; 317 AA.

AC W73507; standard; Protein; 317 AA.

AC W73507; standard; Protein; secreted ligand; 7-Transmembrane receptor;

KW Human Arg-1709 protein; secreted ligand; 7-Transmembrane receptor;

KW heart disease; hypertension; cardiovascular disease; kidney disease;

KW obesity; insuin resistance; diabetes; Central Nervous System disorder;

KW therapy; srRp-1.

SEP 10-3013.

PR 20-301-198; 300313.

PR 20-301-199; 300313

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SSC
                                                                                                                                                                                                                                                                                               !!AA_SEQUENCE 1.0
ID W89910 standard; Protein;
AC W89910;
DT 18-FEB-1999 (first entry)
DE Antigen 1 from cluster 35a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  W73507
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27-APR-1998; U08487.

14-OCT-1997; US-061958.

25-APR-1997; US-045107.

(GENE-) GENELABS TECHNOLOGIES INC.
(GENE-) Fry KE, Lim MY, McAtee CP;
Chow TP, Fry KE, The MY, McAtee CP;
P-009433/01.
                                                                                                                                                                                                                                                                                         The present sequence represents a Helicobacter pylori antigenic protein that is characterised by immunoreactivity with H. pylori-positive antisera. The proteins are highly immunogenic and induce a long-lasting immune response that persists even after antimicriobial treatment. In antibody-detection assays, on sera, plasma, wrine, saliva etc., they are highly sensitive and specific. The specification also describes 69 previously unrecognised immunogenic cluster families. H. pylori antigens infection or to confirm eradication of infection, and in vaccines to protect against H. pylori infection and related diseases (gastritis, peptic uncer, gastric adenocarcinoma/lymphoma).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         analysis.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         peptic ulcer; gastric adenocarcinoma;
Helicobacter pylori.
W09849314-A2.
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New Helicobacter pylori antigens and related nucleic acid sequences

- useful in serological diagnosis and protective vaccines, providing
long-lasting immune response

Claim 16; Page 338; 402pp; English.
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                                                                                                          LSLALKISIK
                                                                                                                                                             IRMANLGEIS
                                                                                                                                    KSAKTTLARI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  r 35a.
family;
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                                                                                                                                                                                                                                                                     2000 15:49
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                                                                                                          DUTDKFLNNM
                                                                                                                                                                                        ADFIINEHPQ
                                                                                                                                                                                                                 INTGGLEYAR
                                                                                                                                                                                                                                          EDTTGEILRH
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                                                                                GVIQTGEEED
                                                                                                                                    ESVDNKLAGA
                                                                                                                                                             PQVVKRVSTV
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                                                                                                                                                                                                                                                                   Type:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   gastritis;
lymphoma.
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                                                                                                                                                              LENKLESLTS
                                                                                                                                                                                        TIALILAHME
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                                                                                                                                                                                                                                                                     Check: 9038
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   diagnosis;
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W89910
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peptic ulcer; gastric adenocarcinoma; ga
Helicobacter pylori.
W09949314-A2.
05-NOV-1998.
27-APR-1998; US-04107.
14-CCT-1997; US-04107.
                                                                                                                                                                  EP-882791-A2.

09-DEC-1998.
303992.
20-MAY-1998; GB-005332.
12-MAY-1997; GB-010755.
23-MAY-1997; GB-010755.
(SMIK ) SMITHKLINE BEECH
Barnes MK, Testa TT;
WPI; 99-011648/02.
N-PSDB; V83106.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           long-lasting immune response claim 1; page 243-244; 402p; English.

Claim 1; page 243-244; 402p; English.

The present sequence represents a Helicobacter pylori antigenic protein that is characterised by immunoreactivity with H. pylori-positive antisera. The proteins are highly immunogenic and induce a long-lasting immune response that persists even after antimicriobial treatment. In antibody-detection assays, on sera, plasma, urine, saliva etc., they are highly sensitive and specific. The specification also describes 69 previously unrecognised immunogenic cluster families. H. pylori antigens are used to detect H. pylori-specific antibodies, for diagnosing infection or to confirm eradication of infection, and in vaccines to protect against H. pylori infection and related diseases (gastritis, peptic ulcer, gastric adenoccarcinoma/lymphoma).
                                                                                                                                                                                                                                                                                                                                        02-MAR 1999 (first entry)
Polypeptide with 98% homology with human Wnt-11.
Mnt-11: disease; detection; cancer; cardiovascular disease; stroke parkinsons disease; schizophrenia; alzheimers disease; acromegaly; vitelliform macular dystrophy; vitreoretinopathy; assay; neurological disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Chow TP, Fry KE, Lim MY, WPI; 99-009433/01.
New Helicobacter pylori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W85590 ST
neurological disease (1918): English. Claim 1; Page 16-17; 21pp; English. This polypeptide has at least 98 % sequence identity with the human which polypeptide has at least 98 % sequence identity with the human whit 1345 makes and sequence. Host cells are used to produce this polypeptide from recombinant constructs which is then used this polypeptide from recombinant constructs which is then used careen for specific agentsts and antagonists (potential therapeutic agents) and to generate Ab (including induction of a protective immune response). Human whit-11 polypeptide is involved in developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      301
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                                                                                                                                 New human Wnt-11 polypeptide - used to diagnose, treat and prevent cancer, cardiovascular disease, developmental disorders, stroke and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (GENE-)
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useful in serological diagnosis and
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;; gastric
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d protective vaccines, providing
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lymphoma.
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New human Wart-11 polypeptide - used to diagnose, treat and prevent PT New human Wart-11 polypeptide - used to diagnose, treat and prevent PT cancer, cardiovascular disease, developmental disorders, stroke and PT neurological disease
PS Claim 17; Page 18-19; 21pp; English.

CC tris polypeptide has at least 95 % sequence identity with the human CC Wart-11 345 amino acid sequence. Host cells are used to produce CC to screen for specific agonists and antagonists (potential therapeutic CC to screen for specific agonists and antagonists (potential therapeutic CC response). Human Wart-11 polypeptide is involved in developmental CC processes and its agonists and antagonists are used for treatment or CC prevention of cancer, cardiovascular disease, stroke, neurological CC disease (e.g. Parkinson's or Alzhahmer's diseases, bipolar/unipolar CC disease (e.g. Parkinson's or Alzhahmer's diseases, bipolar/unipolar CC caromegaly, vitelliform macular dystrophy or vitreoretinopathy.

CC ragments of this polynucleotide sequence can be used for isolating CC related genomic and cDNA clones and for chromosomal or tissue considered consideration (the Wart-11 gene is at chromosome 11q13). The specified CC consesses, or susceptibility to them, are diagnosed by detecting a CC mutation in a genomic whit-11 encoding sequence and/or by analysis for CC presence or amount of Wart-11 (or related mRNA) in a patient sample.
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EP-882791-A2.
09-DEC-1998.
20-MAY-1998; 303992.
12-MAR-1998; GB-005332.
12-MAR-1999; GB-010755.
23-MAY-1997; GB-010755.
(SMIK ) SMITHKLINE BEECHAM PLC.
Barnes MR, Testa TT;
WPT; 99-011648/02.
N-PSDB; V83107.
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W85591 standard; Protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-MAR-1999 (first entry)
Polypeptide with 95% homology with human Wnt-11.
Wnt-11; disease; detection; cancer; cardiovascular disease; stroke; parkinsons disease; schizophrenia; alzheimers disease; acromegaly; vitelliform macular dystrophy; vitreoretinopathy; assay; neurological disease.
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05-SEP-1997
05-SEP-1997
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05-SEP-1997
06-JUN-1997
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W89100 s1
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  US-070923
US-048881
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US-048996
US-0489972
US-049972
US-049972
US-049977644
US-057654
US-057667
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US-048976
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W89100

Length:

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February 11,

2000 15:49

Type: ש

Check: 1342

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R 05-SEP-1997; US-057662.

R 05-SEP-1997; US-057668.

R 05-SEP-1997; US-057668.

R 05-SEP-1997; US-057668.

R 05-SEP-1997; US-057668.

R 05-SEP-1997; US-057765.

R 05-SEP-1997; US-057765.

R 05-SEP-1997; US-057765.

R 06-JUN-1997; US-057765.

R 06-JUN-1997; US-048896.

R 06-JUN-1997; US-048898.

R 06-JUN-1997; US-048898.

R 06-JUN-1997; US-048898.

R 06-JUN-1997; US-048898.

R 06-JUN-1997; US-048995.

R 06-JUN-1997; US-048995.

R 06-JUN-1997; US-0489971.

R 06-JUN-1997; US-0489971.

R 06-JUN-1997; US-0489971.

R 05-SEP-1997; US-0489971.

R 05-SEP-1997; US-057644.

R 05-SEP-1997; US-057764.

R 05-SEP-1997; US-057763.

R 05-SEP-1997; US-057764.

R 05-SEP-1997; US-057766.

R 05-SEP-1997; US
pr New isolated human genes and the secreted polypeptides they encode ruseful for diagnosis and treatment of e.g. cancers neurological pr disorders immune diseases, inflammation or blood disorders limbune diseases, inflammation or blood disorders projected proteins (w86534 to w8756). The secreted protein ce encoding human secreted proteins (w86534 to w8756). The secreted protein ce encoding human secreted with the ATCC under deposit numbers ATCC gene sequences are deposited with the ATCC under deposit numbers ATCC (cc 297974, 97975, 97976, 97977, 209007, 209008, 209011). Host (cc 290011, 20908), 209081, 209082, 209083, 209084, 209085, 209511. Host (cc 290011), 209081, 209082, 209083, 209084, 209085, 209511. Host (cc 290011), 209081, 209082, 209083, 209084, 209085, 209511. Host (cc 290011), 209081, 209082, 209083, 209084, 209085, 209511. Host (cc 290011), 209080, 209081, 209082, 209083, 209084, 209085, 209511. Host (cc 290011), 209080, 209081, 209082, 209083, 209084, 209085, 209511. Host (cc 290011), 209080, 209081, 209082, 209083, 209084, 209085, 209511. Host (cc 290011), 209080, 209083, 209083, 209084, 209085, 209511. Host (cc 290011), 209080, 209083, 209083, 209084, 209085, 209511. Host (cc 290011), 209080, 209083, 209083, 209084, 209085, 209511. Host (cc 290011), 209080, 209083, 209083, 209084, 209085, 209511. Host (cc 290011), 209080, 209083, 209083, 209084, 209085, 209511. Host (cc 290011), 209080, 209083, 209083, 209084, 209085, 209511. Host (cc 290011), 2090802, 209083, 209083, 209084, 209085, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 209010, 
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06-JUN-1997; 0
05-SEP-1997; 0
06-JUN-1997; 0
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Human; secreted protein; fusion protein; gene therapy; protein therapy;
Human; secreted protein; fusion protein; gene therapy; protein therapy;
diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia;
developmental abnormality; foetal deficiency; blood; allergy; renal;
inmune system; asthuma; lymphocytic disease; brain; hepatic; lymphoma;
inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS;
cognitive disorder; schizophrenia; prostate; obesity; osteoclast; thymus;
osteoporosis; arthritis; testis; lung; thyroiditis; thyroid; digestion;
osteoporosis; arthritis; regulation; malabsorption; gastritis; neoplasm.
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WO9854963-A2.
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US-048991
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US-048991
US-048991
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US-057647
US-057647
US-057661
US-057761
US-0577661
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US-0577663
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Caenorhabditis elegans synMuv protein LIN-52. LIN-52; synthetic multivulvar; SynMuv; signal transductio animal model; tumour suppressor; retinoblastoma; cancer; cell proliferation; gene therapy.

signal transduction;

standard; Protein; 161

A

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New isolated human genes and the secreted polypeptides they encode provided for diagnosis and treatment of e.g. cancers, neurological providers, immune diseases, inflammation or blood disorders for claim 11; page 506-507; 772pp; English.

CC thin invention relates to nucleic acid sequences (V84411 to V8463) cc encoding human secreted proteins (W88534 to W88756). The secreted protein (CC gene sequences are deposited with the ATCC under deposit numbers ATCC cc 209011, 209080, 209081, 209082, 209083, 209084, 209085, 209011. Host cells comprising recombinant vectors containing the nucleic acid sequences are used for the recombinant production of the secreted cc proteins. The polypucleotide and amino acid sequences are useful for are useful for preventing, treating or ameliorating medical conditions e.g. cc sequences are mount of the new polypeptides in a sample or by protein or gene therapy. Pathological conditions can be also diagnosed by determining the amount of the new polypeptides in a sample or by colored this was they are most highly expressed in, and include developing cc specific uses are described for each of the polynucleotides, blood cd disorders, developmental abnormalities and foetal deficiencies, blood cd disorders, developmental abnormalities and foetal deficiencies, blood cd disorders, prostate diseases, lymphomas, inflammation, allergies, cischemic shock, Alzheimer's and cognitive disorders involving osteoclasts such as osteoporosis, archititis or malignancies, diseases of testers, lung or restenosis, prostate diseases, obesity, disorders involving osteoclasts such as osteoporosis, archititis or malignancies, diseases of testers, lung or the present sequence represents human secreted protein (see descriptor cc line for gene number and clone identification).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06. JUN. 1997; US-048898.
06. JUN. 1997; US-048898.
06. JUN. 1997; US-048891.
06. JUN. 1997; US-048901.
06. JUN. 1997; US-048901.
06. JUN. 1997; US-048971.
06. JUN. 1997; US-048971.
06. JUN. 1997; US-048971.
06. JUN. 1997; US-048971.
07. SEP-1997; US-057627.
07. SEP-1997; US-057634.
07. SEP-1997; US-057644.
07. SEP-1997; US-057646.
07. SEP-1997; US-057666.
07. SEP-1997; US-057766.
07. SEP-1997; US-057769.
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Brewer LA, Carter KC, Dillon PJ, Ebner R, Endress GA,
Fan P, Feng P, Ferrie AM, Fischer CL, Florence C,
Florence K, Greene JM, Hu J, Kyaw H, Lafleur DW,
Li Y, Moore PA, Ni J, Olsen HS, Rosen CA, Ruben SM,
Shi Y, Soppet DR, Wei Y, Young P, Yu G, Zeng Z;
WPI; 99-059865/05.

NP_SDB; V84445.
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VSPSEYTAAC IGAPGNKPEL February

DEIQPDLREL

METMHRMSHL PPDFEGRQTV SQWLQTLSGM SASDELDDSQ

DKGNLNRCIA SRLLVOYKAA YEEVKLYKNA

DVVSLFITVM

DKLRLEIRAM

Length: 221 MFHGIPATPG

11, 2000 15:49

Type: P

REREKYDNMA

ELFAVVKTMQ Check: 9823

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IIAA_SEQUENCE 1.0

ID W83391 standa.

AC W83391,

DT 29-MAR-1999

DT 29-MAR-1999

Caenorhabditi

KW LIN-52; synth

Each Caenorhabditi

PH 28-MAY-1998;

PH 28-MAY-199;

PH 28-MAY-1998;

PH 28-MAY-199;

PH 28-MAY-1998
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iD w95506 standard; peptide; 446
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This is the amino acid sequence of LIN-52, a novel protein of Caenorhabditis elegans. The lin-52 gene (see V72864) is a novel cancer in mammals. The part of a pathway that may be used as a genetic and biochemical model system for tumour suppression and cancer in mammals. SynMuv pathway genes and proteins may be used to identify genes, which are part of the mammalian pathway and to identify genes, proteins and therapeutic compounds which to identify genes, proteins and therapeutic compounds which condulate this pathway. Pure nucleic acids (see V7289-65) encoding C. elegans SynMuv polypeptides selected from LIN-37, -35, -55, -52, C. 3, -54 and E2F-1 (see W3386-92), are new. Also claimed are:

(1) nucleic acids (see V72866-67) encoding the mouse and human homologues of C. elegans LIN-54; (2) vectors containing the nucleic acids; (3) transgenic cells; (4) a pure mammalian SynMuv family polypeptide, and (5) an antibody which binds to a SynMuv family polypeptide, and treat, especially by gene therapy, conditions cancer involving altered levels of cell proliferation, e.g. SynMuv-
WO9846/US

12-OCT-1998: U07421.

10-APR-1998: US-956985.

24-OCT-1997: US-833610.

A 11-APR-1997: US-834033.

R 11-APR-1997: US-834655.

PR 11-APR-1997: US-834655.

ABBO ) ABBOTT LAB.
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Ceol C, Horvitz HR, Lu X;
WPI: 99-045362/04.
N-PSDB: Y72864.
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28-MAY-1998; U11043.
28-MAY-1997; US-047996.
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Mortierella alpina delta 5 desaturase.
Mortierella alpina delta 5 desaturase.
Delta 5 desaturase; recombinant; fatty acid desaturase; FAD; PUFA; oil;
polyunsaturated fatty acid; linoleic acid; racabidonic acid; linolenic;
polyunsaturated fatty acid; linoleic acid; malnutrition; feeding formula;
stearidonic acid; elcosapentenoic acid; malnutrition; feeding formula;
dietary supplement; prostaglandin; restenosis; angioplasty; inflammation;
rheumatoid arthritis; psoriasis; osteoporosis; cancer; eczema; AIDS;
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Knutzon D,

Leonard

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Mukerji

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Pr plants - useful in human or animal nutrition, as commetties and protection and protection and diabetes provided by the restances of the contains a commetties and diabetes provided by the contains a commetties and diabetes provided by the contains a commetties and contains at least one of the concluding at least one of the concluding at least one of the contains at least one last one last one contains at least one plant cells on the contains and plant cells containing at least one DNA encoding at alpha fatty acid cells containing at least one DNA encoding at another acid desaturated (PDFA). These recombinant cells or plants containing them are used containing them are used containing them are used contained by a containing the
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H. pylori GHPO 1670 protein.

W GHPO protein; Helicobacter infection; gastroduodenal disease; gas

W peptic ulcer disease.

S Helicobacter pylori.

W MO984478-A1.

D 08-OCT-1998; U0-902615.

P 29-JUL-1997; US-902615.

P 101-APR-1997; US-981457.

P 101-APR-1997; US-981277.

A 1-GARAN ) HUMAN GENOME SCI INC.

A (INMR) MERIEUX ORAVAX PASTEUR MERIEUX SERUMS.

A (INMR) MERIEUX ORAVAX FASTEUR MERIEUX SERUMS.

A (INMR) MERIEUX ORAVAX FASTEUR MERIEUX SERUMS.

A (INMR) MERIEUX ORAVAX FASTEUR MERIEUX SERUMS.

N 1-GARAN A, Kleanthous H, Miller C, Oomen RP, Tomb J;

N NPSDB; X14581.

P New isolated Helicobacter polynucleotides - used to develop production of the diagnosis, prevention and treatment of Helicobacter
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        isolated Helicobacter polynucleotides - used to develop products the diagnosis, prevention and treatment of Helicobacter
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GHPO 1670 protes
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ID W98226 standa

standard;

Protein;

115

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31-MAR-1999 (first entry) H. pylori GHPO 1099 protein. GHPO protein; Helicobacter i

infection;

gastroduodenal disease;

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01-APR-1998; U06371.

29-UUL-1997; US-902615.

01-APR-1997; US-833457.

24-JUN-1997; US-881227.
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This sequence represents a Helicobacter pylori GHPO protein of the invention. The polypeptides can be used for preventing or treating the Helicobacter infections, and gastroduodenal diseases associated with these infections, including acute, chronic, and atrophic gastritis, and peptic ulcer diseases, e.g. gastric and duodenal ulcers. They can also be used for the production of antibodies. The products can also be used for detection and diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     31-MAR-1999 (first entry)
H. pylori GHPO 1075 protein.
GHPO protein; Helicobacter infection; gastroduodenal disease; gastritis;
peptic ulcer disease.
Helicobacter pylori.
                                                                                                                                                                                                                                       New isolated Helicobacter polynucleotides - used to develop products for the diagnosis, prevention and treatment of Helicobacter infections and gastrointestinal diseases (claim 8; Page 1616; 2054pp; English.

This sequence represents a Helicobacter pylori GHPO protein of the invention. The polypeptides can be used for preventing or treating inventions, and gastrointeducted his associated with these infections, and gastrointo, and attophic gastritis, and peptic ulcer diseases, e.g. gastric and duodenal diseases. They can also be used for the production of antibodies. The products can also be used for the production of antibodies. The products can also be used for detection and diagnosis.
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(LNMR) MERIEUX ORAVAX PASTEUR MERIEUX SERUMS.
Al-Garawi A. Kleanthous H, Miller C, Oomen RP, Tomb
MPI; 98-542293/46.
51
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                                                                        MIFSFKRTSF
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                                                                    CVSVPVLSIK IFFTLARVSK NNASLNTIKG FLNTPITNAM
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    LMTSTLIHTI
                                                                                                                                February 11,
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    KAFSKLPPKP
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                                                                                                                                         Type: P
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NDLEKLIIQH GLSERVKFFG FVSEKDLPSL ADVKPLNLGY KYLLCVGNRK THKNEKCVIS

RAKRFNWQNV VSEIEMVLTE LTSLTSSLPE VAGDAAILVD

ACDGNK PLSEDAITKG YKGSLGLVFP AFAKADIDPS

ISRLINDSEL SLYEGFGLPV IKLVFTGNPC

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PP 01-MAY-1998; AU0315.

PF 01-MAY-1997; AU-008362.

PR 01-MAY-1997; AU-008365.

PR (UMSY) UNIV SYDNEY.

PA (UMSY) UNIV SYDNEY
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PR (HUMA-) HUMAN GENOME SCI INC.

PA (LIMMA-) HUMAN GENOME SCI INC.

PA (LIMMA-) MERIEUX ORANAX PASTEUR MERIEUX SERUMS.

PA 1-Garawi A, Kleanthous H, Miller C, Oomen RP, Tomb J;

PR WPI; 98-54223/46.

PR New 1solated Helicobacter polynucleotides - used to develop products

PT Net diagnosis, prevention and treatment of Helicobacter

Infections and gastrointestinal diseases

PT Infections and gastrointesting or treating

PT Infection of Including acute, chronic, and atrophic gastricts, and

these infections, Including acute, chronic, and atrophic gastritis, and

PE Infection and diseases, e.g. gastric and duodenal ulcers. They can also be

used for the production of antibodies. The products can also be used for

Sequence 115 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mannosyl transferase involved in O antigen biosynthesis.
O antigen, mannosyl transferase; wbaW gene; diagnosis.
Salmonella enterica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             101
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Helicobacter pylori.
WO9843478-A1.
08-OCT-1998.
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annosyl transferase invol
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W88330

336

February

11,

15:49

Check: 7734

101

FYNFIIKRGC RKAYKIFTVS

NESKERIVAW PLFCSKKFII FIDEISKLLC 2000

SGVNPNKIVT VYNGVSSLFN TIHDLNHLDL NDNSSLFKRI DISKEELYRK CASPLAPLGI Type: P

51

DVVFLPGYIP SWSNKGGIGR

MNNKKVLMDI

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problem in the treating and misging misses and interior mind gagents 32; 251pp; English.

CC Inim 11; Page 232; 251pp; English.

CC moding human secreted proteins (Y01383 to Y01454). The secreted protein gene sequences are deposited with the ATCC under deposit number ATCC conclusions and sequences are used for the recombinant expression of the secreted proteins. The polynucleotide and amino acid sequences are useful for preventing, treating or ameliorating medical conditions e.g. by concein or gene therapy. Pathological conditions can be also diagnosed by concein or gene therapy. Pathological conditions can be also diagnosed by comparing the amount of the new polynucleotides in a sample or by the comparing and triangents, are useful for chromosome identification and mapping; as antisense and triplex-forming therapeutics; in gene therapy; for forensic) identify related sequences or specific mRNA; in preparation conditions and to raise anti-DNA antibodies. Antibodies are useful as immunoassay reagents (including for in vivo imaging) and therapeutically conditions and to raise anti-DNA antibodies. Antibodies are useful as immunoassay reagents (including for in vivo imaging) and therapeutically conditions and to raise anti-DNA antibodies. A very wide range of confidences and triplex forming in polypeptides. A very wide range of inflammation, cancer or other forms of cell proliferation, viral or other infections. The sequences may also be useful in wound healing, to
                                                                                                                                                                                                                                                                                                                                                                                                                                         N-PSDB; x22268.

New nucleic acids encoding secreted human proteins - potentially secreted human proteins - potentially specific useful for treating and diagnosing diseases and identifying specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (HUMA-) HUMAN GENOME SCI INC.
Duan R, Feng P, Ferrie AM, Fl
Duan R, Hu J, Ni J, Rosen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Secreted protein encoded by gene 58 clone HODAZ55. Human; secreted protein; gene therapy; protein therapy; cancer; weight; tumour; chromosome mapping; forensic; haematological disease; allergy; inflammation; cell proliferation; viral infection; wound healing; modulation; appetite; behaviour; food additive; preservative.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .8-MAY-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INC.
M. Florence KA, Fouc.
Rosen CA, Ruben SM, Young
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promoter which is induced in response to a particular operor proteins). Cc described operor proteins of the analyte is analyte (and/or isolating DNA from the protein operating DNA con medium; (c) extracting DNA from these bacteria; (d) identifying mycolic acid bacteria in the vector; (e) cloning vector into suitable host cell; and (f) con medium; (c) extracting DNA from these bacteria; (d) incorporating DNA con medium; (c) extracting DNA from these bacteria; (d) incorporating DNA con medium; (c) extracting DNA from these bacteria; (d) incorporating DNA con medium; (e) extracting DNA from these bacteria; (d) incorporating DNA con medium; (e) extracting DNA from these bacteria; (d) incorporating DNA con medium; (e) extracting DNA from these bacteria; (d) incorporating DNA con medium; (e) extracting DNA for the suitable host cell; and (f) creating DNA con medium; (e) extracting DNA for allows isolation of DNA encoding a component which is induced in response to a particular analyte (and/or promoter which is induced in response to a particular analyte (and/or extraction), which can be used to produce biosensors for the analyte is an environmental pollutant (e.g. from industry or components), especially a hydrophic organic compound such as components of medicine), especially a hydrophic organic compound such as components of concentrations of analytes in samples e.g. for pollution monitoring. The concentrations of analytes in samples e.g. for pollution monitoring. The concentration of also be medicined to produce modified inducible concentrations of analytes and produce modified inducible concentrations of analytes are set and/or operon proteins) whist concentrations of analytes are set and/or operon proteins of the method of inducible enzyme chemistry involved (which may be unknown the present sequence represents a protein from the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Y01440
                                                 W99185 Length: 314
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Sequence 64 AA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE 1.0
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07-JAN-1999.
29-JUN-1998; G01893.
27-JUN-1997; GB-013566.
27-JUN-1997; CAMBRIDGE TECH SERVICES LTD.
(UYCA-) UNIT CAMBRIDGE TECH SERVICES LTD.
Archer JAC, Powell JAC, Roland HJ, Summers
WPI; 99-095760/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rhodococcus corallina ohp operon protein #5.
Rhodococcus corallina; ohp operon; biosensor; mycolic acid bacteria;
inducible promoter; environmental pollutant; industry; medicine.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Isolating DNA encoding inducible promoter from mycolic acid bacteria useful to produce mycolic acid bacterial blosensors for particular analytes, such as environmental pollutants, e.g. from industry or medicine_
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  inducible promoter;
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      MPVALCAMSH
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e 314 AA;
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SPLMGRNDPE QEVIDAVDAA FDHARRFVAD FAPDLIVIFA
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101

MHVDHGFAQA LQLLVGSITA

AQSVGDYGTE

AGPLDVDRDA VPTVPIFINS

AYAVARDVLD

REQRVITAGR DFAAGTAAIQ PLNPEWDRHL LDVLASGDLE

FVGSGGLSHD PPVPQFATAP

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pr Inhibition of virulence genes from Staphylococcus aureus - useful pr for, e.g. screening for potential anti-microbial agents pr for, e.g. screening for potential anti-microbial agents pr for, e.g. screening for potential anti-microbial agents contain 1; Page 179-180; 203pp; English.

CC This is the deduced amino acid sequence of the protein product of a contain 1; Page 179-180; 203pp; English.

CC This is the deduced amino acid sequence of tataphylococcus aureus containing chromosomal insertions containing chromosomal insertions.

CC The nucleotide sequence were identified in a mouse model of bacteraemia. The nucleotide sequence of regions flanking the transposon insertion comparisons were performed to identify these virulence genes (see comparisons were performed to identify these virulence genes (see w97680-724). The plsmy virulence gene product is unknown but comparison and the possible function of their protein products shows 41% identity to the hypothetical 33.7 kpa protein YHCT of Shows 41% identity to the hypothetical 33.7 kpa protein YHCT of ability to interfere with the expression of S. aureus virulence cability to interfere with the expression of S. aureus virulence gene products. Also new is a S. aureus organism containing a gene products. Also new is a S. aureus organism containing a containing and contai
!!AA_SEQUENCE 1.0
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WO9901473-A2.
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Virulence; p15B9; vaccine; antibacterial; antibiotic; screening;
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N-PSDB; X07134.
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WPI; 99-10
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03-JUL-1997; US-887534.
(IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD
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H. pylori ORF hp6p22217_23564012_f1_5
Vaccine; probe; diagnostic; ORF; cell
secreted protein; cellular protein.
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302 AA;
                        ; U19575.
; US-891928.
; US-739150.
; US-759739.
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envelope
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PT New isolated Helicobacter pylori nucleic acids - used to develop PT products for the diagnosis, prevention and treatment of infection by PT products for the diagnosis, prevention and treatment of infection by PT H. pylori and other Helicobacter species
PS Claims 37, 41; Page 224-225; 339pp; English.
CC Recombinant or substantially pure preparations of H. pylori polypeptides CC are disclosed, together with the nucleic acids encoding them. In all, CC cytoplasmic proteins, secreted proteins or other cellular proteins, CC cytoplasmic proteins, secreted proteins or other cellular proteins, CC vaccines containing the nucleic acids or proteins are claimed, as are probes containing at least 8 nucleotides from the nucleic acid sequences. The vaccines are useful for treating or reducing the risk of CC detecting the presence of Helicobacter in a sample. The products are CC detecting the presence of Helicobacter in a sample. The products are CC with the H. pylori life cycle or to inhibit H. pylori infection.
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    Y11062
                                                                                                                                                                                                                                                                                                                                                                                                                                   Helicobacter pylori. W09824475-Al. 11-JUN 1998. 05-DEC-1997; U22104. 14-JUL-1996; US-891928. 05-DEC-1996; US-891928. 25-MAR-1997; US-823745.
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Sequence
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Recombinant or substantially pure preparations of H. pylori polypeptides are disclosed, together with the nucleic acids encoding them. In all, 73 ORFs are shown. The proteins are variously cell envelope proteins, secreted proteins or other cellular proteins. Vaccines containing the nucleic acids or proteins are claimed, as are probes containing at least 8 nucleotides from the nucleic acid sequences. The vaccines are useful for treating or reducing the risk of H. pylori infections, and the probes can be used diagnostically for detecting the presence of Helicobacter in a sample. The products are also of use in screening for compounds having the ability to interfere with the H. pylori life cycle or to inhibit H. pylori infection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    08-JUN-1999 (first entry)
H. pylori ORF 06cp30603_10744075_c3_136 secreted protein.
Vaccine; probe; diagnostic; ORF; cell envelope protein;
vaccine protein; cytoplasmic protein; cellular protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  N-PSDB; x30476. Helicobacter pylori nucleic acids and proteins - used to deproducts for the detection, prevention and treatment of H.
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Alm RA,
WPI; 98
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08-JUN-1999
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Length: 116
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RA, Castriotta LM, Doig
98-333051/29.
DB; X30591.
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February 11,
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Type: P
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W97887

375

February 11, 2000 15:49

Type: ס

Check: 1814

AA,

101

51

LSKLRLETAP

LPKAPPLREL

MPTESDFLMQ NISKDVIRQL

VDGKPKCCFF

KESSKIQYNK

KDGTRYTGIR SLKLDMNPGT

MQKLQLCVYI YLFMLIVAGP VDLNENSEQK ENVEKEGLCN ACTWRQNTKS

ACCCARA CORRES O CONTROL CONTR

14-JUL-1998; IB1197.
15-JAN-1998; US-007761.
14-JUL-1997; US-891789.
(UYLI-) UNIV LIEGE.
Georges M. Grobet L. Pon.
WPI; 99-120869/10.

Increasing muscle mass in mammals

by decreasing myostatin

L, Poncelet

U

Myostatin; human; transforming growth factor beta; double muscling; muscle hyperplasia; transgenic animal.

Human myostatin. W97887; 07-JUN-1999

(first entry)

Properties of the properties of the transforming growth factor-beta superfamily. The invention of the transforming growth factor-beta superfamily. The invention content to the transforming growth factor-beta superfamily. The invention content to the transforming growth factor-beta superfamily. The invention content to the transforming growth factor-beta superfamily. The invention content to the transforming growth factor-beta superfamily. The invention content to the transforming of the myostatin gene and determining composition of the subvine myostatin gene (see X24415-16) has been detected. Certain content to the transforming myostatin content gene are double-muscled. A new method of increasing muscle mass of a mammal having myostatin content substantially complementary to at least a portion of mRNA concoding myostatin and of sufficient length to reduce myostatin encoding myostatin and of sufficient length to reduce myostatin content also be used. Also claimed are: a method for determining muscular content and the mutation; a diagnostic kit for determining the convine; a method for determining the movine; a method for determining the movine; a method for determining the myostatin generally a method for determining the myostatin penotype of an animal; purified myostatin; isolated nucleic acids; a microbial host cell; a probe based on the myostatin movecout animal content and myostatin penotype of an animal having MH phenotype; and a myostatin knockout animal; content animal content and content animal content and content animal content anim

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! AA
21-JAN-1999.
14-JUL-1998;
15-JAN-1998;
14-JUL-1997;
                                                                                                          W97884 standard;
                                     Bovine myostatin.

Myostatin; cattle; bovine; transforming growth factor beta; double muscling; muscle hyperplasia; transgenic animal.

Bos taurus.

W09902667-A1.
                                                                                     W97884;
07-JUN-1999 (first entry)
                                                                                                                                         351
                                                                                                                                        LYFNGKEQII YGKIPAMVVD
                                                                                                          Protein;
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301 251 201

PELEVKVIDI

PKRSRRDFGL VLQNWLKQPE RPVETPTTVF YHATTETIIT

DCDEHSTESR SNLGIEIKAL VQILRLIKPM

CCRYPLTVDF DENGHDLAVT

EAFGWDWIIA

!!AA_SEQUENCE 1.0
ID W97887 standard;

Protein;

375

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101

ATNHODNLSS DIKKIEHEEE

52 Н

DEKVTKEVND

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LMKSKITHFI VISFVLSVLS ACKDEPKKSS QSHQNNTKTT QNNQINQPNK

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TITUD OF THE REAL 
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CC ative myostatin, and a heterologous nucleotide sequence antisense consistency with the antisense sequence.
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14-JUL-1998; US-007761.
15-JAN-1998; US-007761.
14-JUL-1997; US-891789.
(UVLI-) UNIV LIEGE.
Georges M. Grobet L. Poncelet D;
WPI; 99-120869/10.
N-PSDB; XZ4446.
Increasing muscle mass in mammals
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Georges M, Grobet L,
WPI; 99-120869/10.
N-PSDB; X24415, X2446
expression
Disclosure; Page 57; 75pp; English.
This is the amino acid sequence of a mutant bovine myostatin, which is truncated when compared to the wild-type polypeptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUN-1999 (first entry)
Bovine myostatin (mutant form).
Myostatin; cattle; bovine; transforming growth factor beta;
double muscling; muscle hyperplasia; transgenic animal; mutant
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                                                                                                                                                                       decreasing myostatin
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PFLEVKVTDT

PKRSRRDFGL

DCDRISMLSL

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gene are double-muscled. A new method of increasing muscle mass of a mammal having myostatin expressing muscle cells, comprises administration of a nucleic acid molecule substantially complementary to at least a portion of mRNA encoding myostatin and of sufficient length to reduce myostatin expression and thus increase muscle mass. Also claimed are: a method for determining muscle mass also claimed are: a method for determining the genetype of a sample of genetic material; a method for determining muscling in a mammal; a method for determining of a manimal; purified myostatin; isolated nucleic acids; a microbial host cell; a probe based on the myostatin genetype of an animal; purified myostatin; isolated nucleic acids; a microbial host cell; a probe based on the myostatin gene mustatin; and a transgenic bovine having a gene encoding active myostatin and a heterologous nucleotide sequence antisense to that gene, and optionally further containing a gene encoding a secuence 286 AA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               W97884). TI
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ). The mutation results from an 11-bp deletion, i.e. el(11), in the coding region (see X24415) of the myostatin Cattle of the Belgian Blue breed homozygous for the mutant re double-muscled. A new method of increasing musc
286 AA;
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W97885 Length: 286 February 11, 2000 15:49 Туре: Р Check: 1462

101 201 51 GIWQSIDVKT SRLEAIKIQI LSKLRLETAP NISKDAIRQL SSDGSLEDDD MOKLQISVYI YLFMLIVAGP VDLNENSEQK ENVEKEGLCN VLONWLKOPE RPVKTPATVF YHARTETVIT SNLGIEIKAL VQILRLIKPM MPTESDLLTQ LPKAPPLLEL DENGHDLAVT FPEPGEDGLT VEGKPKCCFF KESSKIQYNK IDQFDVQRDA ACLWRENTTS

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Signor began by English.

Signor between the transforming growth factor beta superfamily. The invention relates to factors affecting muscle development in mammals, including the detection of a mutation in the bovine myostatin gene (see x24415-16). Cattle of the Belgian Blue breed homozygous for the mutant gene are double-muscled. A new method of increasing muscle mass of a mammal having myostatin-expressing muscle cells, comprises administration of a nucleic acid molecule substantially comprises administration of a nucleic acid molecule substantially comprises administration of a nucleic acid molecule substantially comprises administration of a nucleic acid molecule muscle cells, comprises administration of a nucleic acid molecule substantially comprises administration of a nucleic acid molecule muscle mass. Comprises a rabsolution of a nucleic acid molecule substantially comprises and thus increase muscle mass. A ribozyme may also be used. Also claimed are: a method for determining muscular hyperplasia (MH) in a mammal using primers based upstream and downstream of the mutation; a diagnostic kit for determining the genotype of a sample of genetic material; a method for
                                                                                                                                                                                                                                                                                                                                             14-JUI-1998; IB1197.
15-AN-1998; US-007761.
14-JUI-1997; US-891789.
(UVII-) UNIV LIEGE.
Georges M. Grobet L, Poncelet D;
WPI; 99-120869/10.
N-PSDB, X24417.
Increasin-
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W97886 stand
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W97-UN-1999 (first entry)
Murine myostatin.
Myostatin; mouse; transfor
double muscling; muscle hy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            mouse; transforming growth cling; muscle hyperplasia; 1
                                                                                                                                                                                                                                                                                                                                                     mammals
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transgenic animal.
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Artificial synthetic trypsin inhibitor gene (WTI-1b) useful for artificial synthetic trypsin inhibitor gene (WTI-1b) useful for insect resistance in transgenic plants - designed on winged bean chemotrypsin inhibitor (WCI-3) using codon and base comparisons bisclosure; page 15; 21pp; English.

This polypeptide comprises chymotrypsin inhibitor WCI-3 of winged bean (Psophocarpus tetragonolobus (L.) DC.). An artificially synthesised gene (see X24282) is claimed which codes for a trypsin inhibitor (see W9783) designed for stable expression in rice. The trypsin inbihitor comprises the winged bean WTI-1B polypeptide having an N-terminal hydrophilic peptide derived from WCI-3 (i.e. amino acid residues 1-24 of WCI-3) utilised for transporting the protein precursor to a vacuole after translation. The synthetic was obtained by: (i) comparing WTI-1B and WCI-3 amino acid sequences, and obtaining homologous amino acid pairs (one each from WTI-1a and b); and (ii) selecting codons from WCI-3 if the amino acid pair is the same, and from WTI-1B if they are different. High usage codons are selected and sequences which cause mRNA stability are modified (see also X24294-96). Also claimed is a plant containing the artificially synthesised gene. Trangenic plants can be produced that are resistant to pests, sequence 207 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                determining MH in a mammal; a method for determining double muscling in a bovine animal; a method for determining the myostatin genotype of an animal; purified myostatin; isolated nucleic acids; a microbial host cell; a probe based on the myostatin gene mutation; transgenic mammals having MH phenotype; and a myostatin knockout animal; and a transgenic bovine having a gene encoding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-JUN-1999 (first entry)
Winged bean chymotrypsin inhibitor wCI-3.
Chymotrypsin inhibitor; wTI-1B;
protease inhibitor; enzyme engineering; protein engineeri
protease inhibitor; enzyme engineering; rice; transge
artificially synthesised gene; winged bean; rice; transge
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07-JUN-
Winged
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W97840 stand
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10-MAR-1999;
01-SEP-1998; 307004.
01-SEP-1997; JP-236332.
(NORQ ) TOHOKU NAT AGRIC
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Artificial
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       standard;
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1. .24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                EXPERIMENT
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e; transgenic
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Length:

207

February

11,

2000

15:49

ID AAC DE DE DE KW

SEQUENCE 1. Y03782 star

standard;

Protein;

AA

11-JUN-1999 (first entry) S. aureus polypeptide. Staphylococcus aureus poly

secretory diarrhoea;

thyroiditis; infective carditis; cerebral abscess; conjunctivitis;

Y03794

Length: 144

February 11, 2000 15:49

Type: P

Check: 7277

51

QPIQRIRDRV

LIASSSISYV

VSQLEDKGWI

TREKDKDDKR VYMACLTEKG RYGLNITEFA VLELLYNKGP

LEQITKEDVK

VLTKDELTIL

QQAFKKLSAQ

MDRTKQSLNV FVGMNRALDT

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PS Claim 31; Page 61-62; 70pp; English.

CC Claim 31; Page 61-62; 70pp; English.

CC Claim 31; Page 61-62; 70pp; English.

CC and the genes (X31851-864) encoding them. Host cells containing vectors comprising the nucleic acid sequences are used for the recombinant CC expression of the proteins. The polypeptides can be used to screen for CC expression of the proteins. The polypeptides can be used to screen for CC entrapy in the proteins and acid sequences are used for treat diseases caused by S. aureus such as thyroiditis, lung abscesses, infective carditis, CC secretory diarrhoea, cerebral abscesses, conjunctivitis, toxic shock Syndrome folliculitis and septic arthritis. Screening for the presence of the polypeptides may be used to diagnose, predict the susceptibility to, CC or stage the progress of these S. aureus diseases and diseases caused by CH elicobacter pylori such as gastric ulcers and gastric adenocarcinoma. There is not much information known about the essential genes expressed by S. aureus during infection but these new polypeptides have been CC by S. aureus during infection but these new polypeptides have been CC addition, these polypeptides can be used to development of antibiotic resistance. The present sequence represents a CC ensures the effectiveness of the compounds in killing S. aureus. In CC ensures the effectiveness of the compounds in killing S. aureus and condition, those polypeptides can be used to effectively diagnose and CC creatiator.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Staphylococcus aureus.

EP-905243-A2.

31-MAR-1999; 306185.

03-AUG-1999; US-055387.

(SMIK ) SMITHKLINE BEECHAM CORP.
(SMIK ) SMITHKLINE BEECHAM PLC.
BUINDAM MKK, LONGETTO MA, WAITEN P.
WPF, 99-192667/17.

New essential polypeptides from St
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Staphylococcus aureus polypeptide; thyroiditis; infective carditis; lung abscess; secretory diarrhoea; cerebral abscess; conjunctivitis toxic shock syndrome; folliculitis; septic arthritis; antibacterial H pylori infection; gastric ulcer; adenocarcinoma.
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Staphylococcus aureus
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  Sequence
                   regulator
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ntial polypeptides from Staphylococcus aureus diseases such as infective endocarditis and
  144 AA;
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pri treating diseases such as infective endocarditis and toxic shock pri treating diseases such as infective endocarditis and toxic shock pri treating diseases such as infective endocarditis and toxic shock pri treating diseases such as infective endocarditis and toxic shock provides new Staphylococcus aureus polypeptides (Y03781-94) and the genes (X31851-864) encoding them. Host cells containing vectors can be invention provides new Staphylococcus are used for the recombinant of the proteins. The polypeptides can be used to screen for ce expression of the proteins. The polypeptides can be used to prevent or treat diseases caused by a natureus such as thyroiditis, lung abscesses, infective carditis, contained to screen for containing the secretory distributis, lung abscesses, infective carditis, contained to syndrome folliculitis and septic arthritis. Screening for the presence of the polypeptides may be used to diseases, infective carditity to, core stage the progress of these S. aureus diseases and diseases caused by G. aureus during infection but these sand gastric adenocarcinoma. CC didntified as essential formation known about the essential genes expressed by S. aureus during infection but these new polypeptides have been contained to development of antibiotic resistance. The present sequence represents a contrained to fantibiotic resistance. The present sequence represents a contrained to a polypeptide which has homology to a B. subtilis probable
    Y03782
                                                                                                                                                                                                                                                                                                                                                                                                                   A_SEQUENCE 1.0
Y02934 standa
Y02934;
21-JAN-1999;
07-UUI-1998;
12-SEP-1997;
12-SEP-1997;
08-UUI-1997;
08-UUI-1997;
08-UUI-1997;
08-UUI-1997;
08-UUI-1997;
08-UUI-1997;
08-UUI-1997;
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O5-AUG-1997; US-055387.
(SMIK ) SMITHKLINE BEECHAM CORP.
(SMIK ) SMITHKLINE BEECHAM PLC.
BURDHAM MKR, LONETTO MA, WARTEN
WPI; 99-192667/17.
                                                                                                                                                                                                                       Fragment of human secreted protein encoded by gene 106.
Fragment of human secreted protein; fusion protein; gene therapy; diagnosis; tissue; cancer; tumour; neurodegenerative disorder; leukaemia; developmental abnormality; foetal deficiency; blood; allergy; renal; immune system; asthma; lymphocytic disease; brain; hepatic; lymphoma; inflammation; ischaemic shock; Alzheimer's disease; restenosis; AIDS; cognitive disorder; schizophrenia; prostate; obseity; osteoclast; thymus; osteoporosis; arthitis; testis; lung; thyroiditis; thyroid; digestion; endocrine; metabolism; regulation; malabsorption; gastrifis; neoplasm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 99-192667,
N-PSDB; X31852.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New essential
                                                                                                                                                                                      Homo sapiens.
WO9902546-A1.
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  U13684.
US-058785.
US-051916.
US-051919.
US-051929.
US-051929.
US-051926.
US-051928.
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Protein; 53
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PT New isolated human genes and the secreted polypeptides they encode principal for diagnosis and treatment of e.g. cancers, neurological principal disorders, immune diseases, inflammation or blood disorders bisclosure; Page 122; 464pp; English.

PS Disclosure; Page 122; 464pp; English of a secreted human protein encoded CC price in the mucleic acid molecule detailed in the descriptor line. The gene cc can be used to generate fusion proteins by linking to the gene to a human cc immunoglobulin Fc portion (e.g. x27302) for increasing the stability of the fused protein as compared to the human protein only.

CC acid sequences: x27311 x2744; amino acid sequences Y02650 Y02788) which care useful for preventing, treating or ameliorating medical conditions can be ce g. by protein or gene therapy. Also, pathological conditions can be composed by determining the amount of the new polypucieotides. CC or by determining the amount of the new polypucieotides. CC which tissues they are most highly expressed in (see X27311 for described cruses).
                        Y02934 Length: 53
                                                                                                                                                                                                                                                                                       Brewer LA, Eb
Olsen HS, Ross
WPI; 99-12077
New isolated
                                                                                                                                                                                                                                                                                                                                          (HUMA-)
                                                   Sequence
                                                                uses)
 Н
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SEP-1997; US-058664.
MA-) HUMAN GENOME SC
WET LA, EDNET R. F1s
en HS, ROSEN CA, Rub
; 99-120770/10.
TLVAGSPCSL
                                                     53 AA;
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US-055723

US-055948

US-055948

US-055940

US-055950

US-055953

US-055953

US-055964

US-055964

US-056366

US-056366
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US-051930.

US-051931.

US-051932.

US-052732.

US-052733.
                        February 11,
 SRWIMAGECH GELVQSDMES QEWERGQVVL SHTSLPWCYV
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hi Y,
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Soppet
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Zeng
                              Check: 2639
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7 Z;
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51 SPR

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II AA_SEQUENCE 1.0

ID Y11970 standard; Protein; 113 AA.

AC Y11970;

DT 18-JUN-1999 (first entry)

DE Human 5 'EST secreted protein SEO ID No: 570.

KW Human; secreted protein; EST; expressed sequence tag; diagnosis;

KW forensic; gene therapy; Chromosome mappling; signal peptide; prostate;

KW upstream regulatory sequence; cytokine activity; cell proliferation;

KW upstream regulatory sequence; cytokine activity; cell proliferation;

KW upstream regulatory sequence; cytokine activity; cell proliferation;

KW upstream regulation; haematopoiesis regulation; tissue growth regulation;

KW thrombolytic; anti-inflammatory; tumour inhibition.

KW thrombolytic; anti-inflammatory; tumour inhibition.

S Homo sapiens.

PN W09905550-A2.

PR 01-AUG-199; US-905144.

PR 01-AUG-199; US-905144.

PA (GEST) GENSET.

Duclett A, Dumas Milne Edwards J, Lacroix B;

DR WPI; 99-153780/13.
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N-PSDB; X40692. New isolated prosta products which may

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PT New polypeptide comprising transcription factor Brn-3a, or its prodefined by the processing nervous system diseases, preventing procellular apoptosis and increasing nervous system diseases, preventing procellular apoptosis and increasing nervous regeneration following procellular apoptosis and increasing nervous regeneration following procession of transcription factor Brn-3a can protect neuronal CC over expression of transcription factor Brn-3a can protect neuronal CC of the BC1-2 gene in neuronal cells and this activates expression CC of the BC1-2 gene. Both the anti-apoptopic effect of Brn-3a and its ability to activate expression of BC1-2 are mediated by the N-terminal CC of the BC1-2 gene. Both the anti-apoptopic effect of Brn-3a and its ability to activate expression of BC1-2 are mediated by the N-terminal coles in the regulation of selective apoptosis during development of the neurons system. The stimulation of Bc1-2 expression by Brn-3a in a neuron specific manner and consequent protection of neuronal cells from apoptosis suggests that Brn-3a my co-ordinate some aspects of neuronal reorganisation during development or following injury. The cleavest of Brn-3a expression by either pharmacological means of comparision of the protection of protection of Brn-3a expression by either pharmacological means.
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WO9905272-A1.
04-FEB-1999.
27-JUL-1998; G02228.
10-DEC-1997; US-988476.
25-JUL-1997; GB-018823.
25-JUL-1997; GB-018823.
(UNLO) UNIV COLLEGE LONDON.
elevation of (compositions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Brn-3a polypeptide.

Brn-3a; Bcl-2; neurons; neuronal cells; apoptosis; cell death; CNS; Brn-3a; Bcl-2; neurons; neuronal cells; apoptosis; cell death; CNS; PNS; central nervous system; parasympathetic nervous system; development; injury; neurotrophic factor; nerve growth factor; NGF; celliary neurotrophic factor; CNTF; brain derived neurotrophic factor; BNTF; neurotrophin; NT-3; NT-4; NT-5; neurodegenerative disease; familial dysautonomia; infantile muscular dystrophy; Parkinson's disease; Alzheimer's disease.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Latchman DS, Smith WPI; 99-142928/12. N-PSDB; X09010.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  standard; Protein; 423
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PT New isolated prostate-derived nucleic acids - used to develop products which may have cytokine, immune regulatory, haematopoiesis PT regulating, anti-inflammatory or tumour inhibition activity atti-inflammatory or tumour inhibition activity atti-inflammatory or tumour inhibition activity (CC x40438 to x40715 represent 5' expressed sequence tags (ESTs) for human CC x11716 to Y11993 respectively. The proteins given represent the signal CC peptide and an N-terminal fragment of a secreted protein. The nucleic CC acid sequences can be used for producing secreted human gene products. The proteins obtained may have cytokine activity, cell proliferation and CC differentiation activity, haematopoiesis regulating activity, tissue CC growth regulating activity, reproductive hormore regulating activity, componently and activity, anti-inflammatory activity, tumour inhibition CC activity or other activities. The products can be used in forensic, gene CC therapy and chromosome mapping promoters on be used in forensic, gene CC encoding the signal peptides can be used for obtaining corresponding promoters sequences. The nucleic acids centration of a polypeptide or the insertion of a polypeptide into a cell. Sequence 113 AA;
                                                                                                                                                                                                                                                                                                                                                                                    888888888
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                                                         201
                                                                                                            151
                                                                                                                                                                  101
                                                                                                                                                                                                                                                                                                                                                                                                                              neurotrophic factors, nerve growth factor (NGF), ciliary neurotrophic factor (CNTF), brain-derived neurotrophic factor (BNTF), and neurotrophins NT-3 and NT-4/5) or gene therapy may represent a method for treating human diseases associated with excessive neuronal cell death and/or lack of nerve regeneration, especially neurodegenerative diseases such as familial dysautonomia and infantile muscular dystrophy, and Parkinson's and Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                    disease
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401 351 301 251

NVVRVWFCNQ ROKOKRMKFS

ATY

PELFNGGEKK

RKRTSIAAPE KRSLEAYFAV

STICRFESLT SICDSDTDPR

LSHNNMIALK ELEAFAERFK

PILQAWLEEA EGAQREKMNK QPRPSSEKIA AIAEKLDLKK

QRRIKLGVTQ

ADVGSALANI

AAVVGAAGLA

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THE SERVICE SERVICE SERVICES S
                                                                                                 DR WPI; 99-244657/21.

PT Use of an MHC class II-associated invariant chain protein (CLIP) for PT treating or preventing an immune disorder.

CC Theis sequence represents the human CLIP immunomodulatory peptide.

CC Theinvention relates to a method of treating or preventing an immune contains requested invariant to chain protein (CLIP) to an affected/susceptible mammal. The CLIP peptides are useful as immunogenic compositions for: (1) increasing the level of CLIP cells; (2) decreasing the level of TH1 cells; (3) preventing or cell contain an immune response; (4) reducing or preventing are response to an antigen; (5) reducing or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (5) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen presentation by APCs; and (6) treating or preventing antigen prev
for screening to arrive for ability to arrive for a finite feron-gamma.
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WPI; 99
Use of
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CLIP; immunomodulatory peptide; immune disorder; TH1 cell reduction; MHC class II-associated invariant chain protein; transplant rejection; FH2 cell; immune response; T cell response; T cell activation; arthritis; autoimmune disorder; autoimmune diabetes; multiple sclerosis; systemic lupus erythematosus; cytokine production; human.
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16-NOV-1998.

16-MAY-1997, 205680.

16-MAY-1997; CA-205680.

(UYWO-) UNIV WESTERN ONTARIO.
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Y05204 $\boldsymbol{\mu}$ Length: 24 February 11, 2000 15:49 Type: ש 3125

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THE REPORT OF THE RESERVE OF THE RES
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        Human 5' EST secreted protein SEQ ID No: 593.

Human, secreted protein; EST; expressed sequence tag; diagnosis; forensic; gene therapy; chromosome mapping; signal peptide; prostate; upstream regulatory sequence; cytokine activity; cell proliferation; differentiation; haematopolesis regulation; tissue growth regulation; reproductive hormone regulation; chemotactic; chemokinetic; haemostatic; thrombolytic; anti-inflammatory; tumour inhibition.
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16-SEP-1997; GB-019727.
(RHON) RHONE-POULENC AGRIC L
COLE DJ, Cummins I, Edwards R
WPI; 99-244035/20.
N-PSDB; X25152.
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DUCLET A, Dumas Milne Edwards J, Lacroix B;

WPI, 99-153780/13.

N-PSDB; X40715.
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WO9906550-A2.
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31-JUL-1998; IB1232.
01-AUG-1997; US-905144.
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Y05545 standard;
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New isolated glutathione transferase subunit polynucleotides Claim 3; page 100; 101pp; English.

The present sequence represents ICP and ICV, wheat glutathione transferase (GST) subunits that resemble the type III GSTs of maize. ICP/ICV cDNA (see x25152) was isolated from a cDNA library prepared from fenchlorazole-ethyl (herbicide safener) treated wheat shoots.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-701-1999 (first entry)
Wheat Type III glutathione transferase subunit ICP/ICV.
Glutathione transferase; 6507; glutathione peroxidase; wheat;
ICV; herbicide resistance; transgenic plant.
Triticum aestivum.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             51
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Y07041 Length: 207

MSPLLRRLLL

February :

11,

2000 15:49

Type:

Check: 1679

MGTVAKQLVP LGEMSLEEHS

APVSQPDAPG SCVTVQRCGG

HQRKVVSWID

PTGQHQVRMQ

51 101

EVVVPLTVEL ILMIRYPSSQ

QCECRPKKKD SAVKPDRAAT

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88888888888
             ID Y07041 st
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                                             New isolated cancer associated nucleic acids and polypeptides - prisolated using sera from cancer patients, used to develop products prisolated using sera from cancer patients, used to develop products prisolated using sera from cancer patients, used to develop products prisolated using sera from cancer associated ancers prisolated repairs and in the page 426-427; 787pp; English.

So bisclosure; page 426-427; 787pp; English.

Co bisclosure; page 426-427; 787pp; English.

Co bisclosure; page 426-427; 787pp; English.

Co product complexes of human cancer associated antigen precursor coded for by expression of a human cancer associated antigen precursor coded for by expression product with an agent that specifically bisclosure; page 426-427; 787pp; English and Contacting a content of the NAM, an expression product or a fragment of an expression of product complexed with an HLA molecule; and (b) determining the content of the disorder. The products and methods can be used in the diagnosis, monitoring, research, or treatment of conditions or the diagnosis, monitoring, research, or treatment of conditions or the invention provides nucleic acid sequences and encoded polypeptides of the invention provides nucleic acid sequences and encoded polypeptides which are cancer associated antigen precursors expressed in human breast cancer, renal cancer, colon cancer, gastric cancer, prostate cancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          15-UL-1998; U14679.
22-UNN-1998; US-102322.
17-UNL-1997; US-896164.
10-OCT-1997; US-061599.
10-OCT-1997; US-061599.
10-OCT-1997; US-061765.
10-OCT-1997; US-041765.
11-OCT-1997; US-041697.
(LUDW-) LUDWIG INST CANCER RES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention provides wheat GST subunits (see Y05537-45) active in herbicide metabolism. This is fundamental to understanding GST detoxification in wheat and in the development of transgenic herbicide resistant plants expressing wheat GSTs. The invention herbicide resistant plants expressing wheat GSTs. The invention provides methods of identifying compounds capable of metabolism by GST, or compounds that induce GST expression in graminaceous plants, and for determining the GST level in a sample of seed or flour. Transgenic plants, host cells used for production of GST subunits and GST dimeric proteins, and vectors are also provided.

Sequence 243 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             101
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Breast cancer associated antigen precursor sequence.
Cancer associated antigen; diagnosis; research; treatment;
Cancer associated antigen; diagnosis; research; treatment;
breast cancer; colon cancer; gastric cancer; renal cancer;
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j cancer.
207 AA;
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DR WPI; 99-132448/11.

PT New isolated cancer associated nucleic acids and polypeptides - isolated using sera from cancer patients, used to develop products PT isolated using sera from cancer patients, used to develop products PT isolated using sera from cancer patients, used to develop products PT isolated using sera from cancer isolated cancers

PT Disclosure; Page 391-392; 787pp; English.

CC The invention relates to a method for diagnosing a disorder characterised by expression of a human cancer associated antigen precursor coded for by a cc nucleic acid molecule (NAM). The method comprises: (a) contacting a cc biological sample isolated from a subject with an agent that specifically binds to the NAM, an expression product or a fragment of an expression cc interaction between the agent and the NAM or the expression product as a codetermination of the disorder. The products and methods can be used in the diagnosis, monitoring, research, or treatment of conditions characterised by the expression of various cancer associated antigens. The invention provides nucleic acid sequences and encoded polypeptides which are cancer associated antigen precursors expressed in human breast cancer, renal cancer, colon cancer, gastric cancer, prostate cancer and
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22-UUR-1998; US-102322.
17-JUL-1997; US-896164.
10-OCT-1997; US-061599.
10-OCT-1997; US-061765.
10-OCT-1997; US-061765.
10-OCT-1997; GB-021697.
11-OCT-1997; GB-021697.
(LUDW-) LUDWIG INST CANCER RES.
(LUDW-) LUDWIG INST CANCER RES.
Chen Y, Gout I, Gure A, OHare M, Oh
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W09904265-A2.
28-JAN-1999.
                                                        Human regulatory molecule, HRM-9 protein.

Human regulatory molecule, HRM-9; cytostatic activity; immune modulator; transcription factor; enhancer; cell proliferation stimulation; cancer; treatment; microarray; detection; diagnosis; cell proliferation disease; leukemia; lymphoma; myeloma; adenocarcinoma; sarcoma; bladder; bone; leukemia; lymphoma; myeloma; adenocarcinoma; sarcoma; bladder; bone; brain; lung; liver; ovary; skin; teratocarcinoma; fine process; alicenteriations; central disease;
                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE 1.0
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myasthenia gravis.
                                allergy; asthma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                 v93953 standard;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LHMTCAKCNH
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                                    diabetes; multiple
                                                                                                                                                                                                                                                                                                                                                                 Protein;
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MJ, Stockert
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nal cancer; lung cancer;
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POS XXX XXX DD DAC

February 11,

.2000 15:49

Type:

Check:

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II AA_SEQUENCE 1.0

ID Y04959
AC Y04959
DT 06-JUL-1999 (first entry)
DE Mycobacterium species protein
KW Secreted protein; Mycobacter
KW Mycobacterium sp.
PN W9999186-A2.
PN W9099186-A2.
PN W9099186-A2.
PN W9099186-A2.
PN W1-SEB-1997; FR-010404
PN (INSP) INST PASTEUR.
PN Gicquel B, Lim EM, Pelicic \
PN Gicquel B, Lim EM, Pelicic \
PN Gicquel B, Lim EM, Pelicic \
PN WPI; 99-181045/15.
PN WPI; 99-181045/15.
DR WPI; 99-181045/15.
DR WPI; 99-181045/15.
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DR WPI; 99-181045/15.
PN WPI; 99-181045/15.
DR WPI; 99-181045/15.
DR WPI; 99-181045/15.
DR WPI; 99-181045/15.
CR identifying coding or promoi infection-associated protein for infection associated protein for infection asso
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         Mycobacterial DNA vectors containing reporter constructs - for identifying coding or promoter sequences involved in infection-associated protein expression claim 32; Fig 42D; 309pp; French.

Claim 32; Fig 42D; 309pp; French.

Sequences Y04742-Y05000 and Y07201-Y07204 represent secreted proteins from various Mycobacterium species microorganisms. The encoding nucleotide sequences can be used as primers and probes for methods for detecting and identifying mycobacteria, especially belonging to the M. tuberculosis complex. The encoded proteins can be used in vaccines for immunisation against a bacterial or viral infection. Sequence 273 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes novel human regulatory molecules (HRM) which have cytostatic activity and act as immune modulators, transcription factors or enhancers. The HRMs can be used to stimulate cell proliferation. Antagonists and agonists of the proteins of the invention can be used to treat cancer. The encoding nucleic acids can be used in microarrays to detect polynucleotides (and their expression levels) that encode HRMs in a biological sample. The HRMs and microarrays can be used diagnose, treat or prevent cell proliferation diseases especially cancer, and the province of the content of the con
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Mycobacterium species protein sequence 42D. Secreted protein; Mycobacterium; primer; PCR; amplification; hybridisation; detection; vaccine; immunisation; infection.
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Shah P, Yue H;
WPI; 99-254710/21.
N-PSDB; X24067.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-SEP-1998; U19839.
23-SEP-1997; US-933750
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14-AUG-1998; F01813. 11-SEP-1997; FR-011325. 14-AUG-1997; FR-010404. (INSP) INST PASTECR. Gicquel B, Lim EM, Pelic: Mycobacterial DNA vectors containing reporter constructs - for identifying coding or promoter sequences involved in infection-associated protein expression claim 32; Fig 42F; 309pp; French.
Sequences Y04742-Y05000 and Y07201-Y07204 represent secreted proteins from various Mycobacterium species microorganisms. The encoding nucleotide sequences can be used as primers and probes for methods for detecting and identifying mycobacteria, especially belonging to the M. tuberculosis complex. The encoded proteins can be used in vaccines for immunisation against a bacterial or viral infection. Mycobacterium species protein sequence 42F. Secreted protein; Mycobacterium; primer; po Secreted protein; detection; vaccine; immunisa SEQUENCE 1. 251 151 201 101 Juigueno 51 cobacterium sp. eno A; 99-181045/15. RYAIHELALV MANKRGNAGQ PLPLSDRDDD HMQGHWLLAR LGKRVLRPGG LVVEHVVTAS GDVRVTDAAD AEVTDADVLE LAPGLGRTAA standard; Lim EM, Pelicic V, (first entry) AVAIVAHKPH MALLQPRRVI PDDVAEQVRT TGLSDASADV Protein; 280 VDS VIGEAMLIMO ADEGLIGALR DLRQSLARAL EILARNPRSY Portnoi D, Ą immer; PCR; amplification; immunisation; infection. FAGNLLIHRA ARRRVLLMRH GNAAKHTIVA VGAESDPNAA KVNARPLTVA Goguet PCR; de EWSHLLAGHG EAARVLRPGG NLVRHVLAGR VELTRILLAR la Salmoniere probe;

Y04960 Length: 280 February 11, 2000 15:49 Type: P Check: 8320

51 TRTLLARAEV TDADVLELAP GLGRTAAEIL ARNPRSYVGA ESDPNAANLV

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KPAEVGSMAN

KRGNAGQPLP

LSDRDDDHMQ GHWLLARLGK RVLRPGGVEL

RHVLAGRGDV RVTDAADTGL SDASADVVIG EAMLIMOGNA AKHTIVAEAA

RVLRPGGRYA IHELALVPDD VAEQVRIDLR QSLARALKVN ARPLTVAEWS

201 HLLAGHGLVV EHVVTASMAL LOPRRVIADE GLLGALRFAG NLLIHRAARR

251 RVLLMRHTFR RHRERLTAVA IVAHKPHVDS

IIIAA AC DI DE DI DE CON DE COND. 25-FEB-1999. 14-AUG-1998; F01813. 11-SEP-1997; FR-011325. 14-AUG-1997; FR-010404. (INSP) INST PASTEUR. SEQUENCE 1.0 Guigueno A; WPI; 99-181045/15 Gicquel B, 06-JUL-1999 (first entry)
Mycobacterium species protein sequence 31C.
Secreted protein; Mycobacterium; primer; PC
hybridisation; detection; vaccine; immunisa Mycobacterium sp. standard; Lim EM, Protein; Pelicic V, 72 ₽ Portnoi D, immunisation; Goguet PCR; amplification; probe;
on; infection. de la Salmoniere М

nflammation.

Homo sapiens. EP-906953-A1.

EP-905222 07-APR-1999. 16-SEP-1997; 116061. (16-SEP-1997; EP-116061. (1STF) ARS APPLIED RES SYSTEMS HOLDING NV

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!!AA_SEQUENCE 1.0
ID Y07239 standard; Protein; 141 AA.

101

PRQFAGYLSQ GOPIAFDAYS MAGDQELELR

51

VAQLFGDVTG FDVPLYTLAE

ARVAGVQPQR ASRYLVVPRA

HHIRPVRLRG PLGGVGCLRH TLATWADGYE RRPANAPAVQ

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06-JUL-1999 (first entry)
Fragment of human placental STAT3 protein.
Placenta: isoform, human, STAT3; intracellular; transcription factor;
Signal Transducer and Activator of Transcription; allele; growth arrestic acute-phase protein; monocytic cell; myeloma; autoimmune dise

Y04743 Length: 110

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Y04901 11-SEP-1997; 1 14-AUG-1997; 1 (INSP) INST 1 Gicquel B, Lir Mycobacterial DNA vectors containing reporter constructs - for identifying coding or promoter sequences involved in infection-associated protein expression claim 32; Fig 1, 309pp; French, 197201-197204 represent secreted proteins sequences v04742-v05000 and v07201-v07204 represent secreted proteins from various Mycobacterium species microorganisms. The encoding nucleotide sequences can be used as primers and probes for methods for detecting and identifying mycobacteria, especially belonging to the M. tuberculosis complex. The encoded proteins can be used in vaccines for immunisation against a bacterial or viral infection. Guigueno A; WPI; 99-181045/15. Mycobacterial DNA vectors containing reporter constructs - for identifying coding or promoter sequences involved in infection-associated protein expression claim 32; Fig 31C; 309pp; French.

Sequences Y04742-Y05000 and Y07201-Y07204 represent secreted proteins from various Mycobacterium species microorganisms. The encoding nucleotide sequences can be used as primers and probes for methods for detecting and identifying mycobacteria, especially belonging to the M. tuberculosis complex. The encoded proteins can be used in vaccines for immunisation against a bacterial or viral infection. Sequence N-PSDB; SEQUENCE 1.0 Sequence secreted protein; hybridisation; det N-PSDB; X34154 fycobacterium)6-JUL-1999 51 04743 standard; ycobacterium sp. Ь Length: 72 February 11, ANTREGSATA PGSISRATRR SAASISARRA for immunisation against a
110 AA; for immunisation against
72 AA; Lim FR-011325. FR-010404. (first entry) PASTEUR. speci EM, species protein sequence 1#2.
in; Mycobacterium; primer; PCR;
detection; vaccine; immunisatio ALPAPWLLNN TSQPRTLPVC GRINDPNSTN TRIATTKALA Protein; Pelicic 110 AA ۲, SR 2000 15:49 Portnoi D, immunisation; Goguet Type: P amplification; probe; фe Check: 6286 la Salmoniere

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                                                                                                                                                                                                                                                                                                                                                                                           New allelic variant of human STAT3 useful in treating autoimmune or inflammatory diseases Claim 2: Page 10-11; 22pp; English.

This sequence represents a fragment of the hepatic allelic isoform of human Signal Transducer and Activator of Transcription (STAT3) intracellular transcription factor (Akira et al., Cell 77, 63-71 (1994)). The invention relates to isolation of allelic variants of the placental harras sequence harmy plays a role in the upregulation of hepatic acute phase proteins, growth arrest of monocytic cells and in the survival of myeloma cells and so may be used to treat or diagnose autoimmune or inflammatory diseases.

Sequence 141 AA;
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EP-906953-A1.
07-APR-1999.
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Disclosure; Page 8-9; 2pp; English.
Disclosure; Page 8-9; 2pp; English.
This sequence represents a fragment of the placental isoform of human Signal Transducer and Activator of Transcription (STAT3) intracellular transcription factor (Akira et al., Cell 77, 63-71 (1994)). The invention relates to isolation of allelic variants of the placental hSTAT3 sequence. hSTAT3 plays a role in the upregulation of hepatic acute-phase proteins, growth arrest of monocytic cells and in the survival of myeloma cells and so may be used to treat or diagnose autoimmune or inflammatory diseases.

Sequence 141 AA;
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Y07240 standard; Protein; 141 AA.
                                            06-JUL-1999 (first entry)
Fragment of mouse hepatic STAT3 protein.
Placenta; isoform; human; STAT3; intracellular; transcription factor;
Signal Transducer and Activator of Transcription; allele; growth arrest;
hepatic acute-phase protein; monocytic cell; myeloma; autoimmune disease;
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16-SEP-1997; EP-116061.
(ISTF ) ARS APPLIED RES SYSTEMS HOLDING
Della Pietra L, Serlupi-Crescenzi O;
WPI; 99-207107/18.
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Fragment of human hepatic STAT3 protein.
Placenta; isoform; human; STAT3; intracellular; transcription factor;
Signal Transducer and Activator of Transcription; allele; growth arrest;
hepatic acute-phase protein; monocytic cell; myeloma; autoimmune disease;
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DB; X29975.
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                                                                                                                                                                                                                                                                                                                       WLDNIIDLVK KYILALWNEG YIMGFISKER
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                                                                                                                                                                             standard; Protein; 141 AA.
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PD 07-APR-1999.

PF 16-SEP-1997; 116061.

PR (LSTF ) ARS APPLIED RES SYSTEMS HOLDING NV.

PR (LSTF ) ARS APPLIED RES SYSTEMS HOLDING NV.

PR Della Pietra L, Serlupi-Crescenzi O;

PR NPSDB; X29977.

DR N-PSDB; X29977.

PR New Allelic variant of human STAT3 useful in treating autoimmune or prinflammatory diseases

PI sclosure; Page 12-13; 22pp; English.

CC of mouse Signal Transducer and Activator of Transcription (STAT3)

CC of mouse Signal Transducer and Activator of Transcription (STAT3)

CC of nouse Signal Transducer and Activator of Transcription (STAT3)

CC of nouse Signal Transducer and Activator of Transcription (STAT3)

CC of the placental human STAT3 sequence. hSTAT3 plays a role in the placental human STAT3 sequence. hSTAT3 plays a role in the ccells and in the survival of myeloma cells and so may be used to treat CC or diagnose autoimmune or inflammatory diseases.

SQ Sequence 141 AA;
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Y07241 Length: 141 February 11, 2000 15:49 Type: P Check: 3064

51 WLDNIIDLVK KYILALWNEG IQSVEPYTKQ QLNNMSFAEI IMGYKIMDAT YIMGFISKER ERAILSTKPP GTFLLRFSES

101 NILVSPLVYL YPDIPKEEAF GKYCRPESQE SKEGGVTFTW VEKDISGKTQ HPEADPGSSA